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                                                                                                                                                                                Fungal detection; bacterial detection; probe; primer; ss.
                                                                                                                                                                                                           Aspergillus auricomus probe.
                                                                                                                                                                                                                                                                          ABK29246 standard; DNA; 22
                                                         Sn (ISSn)
                                                                                                                                                                                                                                      23-APR-2002
                                                                                                                                                                                                                                                           ABK29246;
                                                                                                                                                            spergillus auricomus.
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AAV61691
AAV15500
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AAQ57362
AAT18458
AAT08056
AAT85301
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Primer 467R for R
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Alzheimer's diseas
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                                                                                                                    Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe
                                                                                              Claim 87; Page
                                                                                                                                                                          WPI; 2002-098078/13.
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25;

110pp; English.

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RESULT 2
ABK29280/c
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Best Local
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  Haugland R,
                                 (USSI ) US ENVIRONMENTAL PROTECTION AGENCY.
                                                                                                                   13-JUN-2001; 2001WO-US18892
                                                                               13-JUN-2000; 2000US-0593012
                                                                                                                                                                                                       WO200196612-A2
                                                                                                                                                                                                                                               Aspergillus sclerotiorum
                                                                                                                                                                                                                                                                              Fungal detection; bacterial detection;
                                                                                                                                                                                                                                                                                                                      Aspergillus sclerotiorum probe.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of detecting and quantifying fungi a bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and the continuous continuous chain reaction (PCR) and fluorescent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting and quantifying fungi and bacteria, involve sequence of the fungus, extracting the DNA from the subjecting the DNA to polymerase chain reaction and
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4.9e+02;
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                                                                                                                                                                                                                                                                      The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the bapple, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific partice sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and fluorescent processing the DNA to polymerase chain reaction and the DNA to polymerase chain reaction reaction and the DNA to polymerase chain reaction reaction reaction reaction reaction reaction react
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ilarity 100.0%;
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RESULT

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RESULT 5
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                       WO200196612-A2
                                                                        Fungal detection; bacterial
                                                                                                Aspergillus niveus probe.
                                                                                                                           23-APR-2002
                                                                                                                                                     ABK29265;
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                                               Aspergillus niveus.
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                                                                          detection; probe;
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                                                                                                                                                                                                                                                                                                               1.7e+03;
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                                                                          primer; ss
                                                                                                                                                                                                                                                                                                                          Length 16;
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RESULT 6
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Best Local :
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                        Detecting and quantifying fungi and bacteria, involves obtaining sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16
                                                                                                                                                                                                           Haugland R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aspergillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fungal detection; bacterial detection; probe; primer; ss
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ilarity 100.0%;
Conservative (
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fluorescent probe
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Best Local S
Matches 14
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                          The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
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                                                                                                                                                                                                                                                           Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent pro
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Pred. No. 2.
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Query Match

Sequence

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BP; 3 A; 7 C; 8 G; 5 T; 0 other;

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Score 12.8;

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                                                      Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                sequences unique to the species/group of species but common to all isolates of the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe sequences (ABN99429-ABN99653) for detection of specified fungifungal groups or specified bacteria. The method enables detection and quantification of specific fungi or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment of fungi or bacteria associated with health problems such as infections, gastroenteritis, ulcers, asthma, allergies and sinusitis. It is also useful to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary haemorrhage from levels of Stachybotrys chartarum in air samples in
                                                                                                                                                                                                                                                                                                                         The invention relates to specific fungi or bacteria detected and quantified in a sample by extracting and recovering DNA from the organism in the sample and hybridising and amplifying the DNA sequence using
                                                                                                                                                                                                                                                                                                                                                                                  Claim 65;
                                                                                                                                                                                                                                                                                                                                                                                                                          Detection and quantification of specific fungi or bacteria useful e.g. in medical diagnosis and treatment of fungal and bacterial conditions, by hybridizing and amplifying DNA using sequences unique to the specie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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                                                                                                           Sequence 15
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14-APR-1999;
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 TTGCCTCGGCGGGA 14
                          TTGCTTCGGCGGGA 15
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allergy; sinusitis; PCR; primer; ss.
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                                                                  Score 12.4; DB 2
Pred. No. 3.5e+03
                                                         Mismatches
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ABK29220
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fungal detection; bacterial detection; probe; primer; ss
                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of detecting and quantifying bacteria, involving obtaining a sequence of the microorganism t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 65; Page 21; 110pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-098078/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Haugland R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUN-2000; 2000US-0593012
                                      Penicillium expansum
                                                               Fungal detection;
                                                                                          Penicillium expansum reverse PCR primer.
                                                                                                                     23-APR-2002
                                                                                                                                              ABK29366;
                                                                                                                                                                         ABK29366 standard; DNA; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  analysis
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                                                                                                                                                                                                                                                                                             13;
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                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                  15 BP; 1 A; 4 C; 6 G; 4 T; 0 other;
                                                                                                                                                                                                                                        TIGCCICGGCGGGA 14
                                                                                                                                                                                                                                                                   TIGCTICGGCGGGA 15
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                                                                                                                     (first entry)
                                                                 bacterial
                                                                                                                                                                                                                                                                                                         72.9%;
                                                                                                                                                                                                                                                                                             0
                                                                 detection; probe;
                                                                                                                                                                                                                                                                                                           Score 12.4; DB 24
Pred. No. 3.5e+03;
                                                                                                                                                                                                                                                                                                Mismatches
                                                                  primer;
                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                Indels
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent pro
                                                                                                                                                                                                                                                                                                                                                                    ABN99619 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16 BP; 3 A; 7 C; 5 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 116; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-098078/13
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                                                                                                                                                                                        US6387652-B1
                                                                                                                                                                                                                                                Pungus; bacteria; quantification; detection; infection;
ulcer; asthma; allergy; sinusitis; probe; ss.
                                                                                                                                                                                                                                                                                         Fungi probe
                                                                                                                                                                                                                                                                                                                    05-AUG-2002
                                                                                                                                                                                                                                                                                                                                             ABN99619;
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Detection and quantification of specific
                                                                                                 15-APR-1998;
14-APR-1999;
                                                                                                                                                                 14-MAY-2002
                                                                                                                                                                                                                            Trichoderma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (USSI ) US
                         WPI; 2002-462353/49
                                                                                                                                      13-JUN-2000; 2000US-0593012
                                                                         (USSI ) US ENVIRONMENTAL PROTECTION AGENCY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     GTAGCTTCGGCGGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTTGCTTCGGCGGG 14
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                                                                                                                                                                                                                asperellum.
                                                                                                                                                                                                                                                                                         SEQ ID NO 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vesper SJ;
                                                Vesper
                                                                                                                                                                                                                                                                                                                   (first
                                                                                                  98US-081773P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31, 110pp;
                                                                                                                                                                                                                                                                                                                                                                     DNA;
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Pred. No. 3.5e+03;
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fungi or bacteria useful
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                                                                                                                                                                                                                                                                 gastroenteritis;
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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences unique to the species/group of species but common to all isolates of the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe sequences (ABN99429-ABN99653) for detection of specified fungi/fungal groups or specified bacteria. The method enables detection and quantification of specific fungi or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment of fungi or bacteria associated with health problems such as infections, gastroenterits, ulcers, asthma, allergies and sinusitis. It is also useful to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary haemorrhage from levels of Stachybotrys chartarum in air samples in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in medical diagnosis and treatment of fungal and bacterial conditions, by hybridizing and amplifying DNA using sequences unique to the specie or group of species -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to specific fungi or bacteria detected and quantified in a sample by extracting and recovering DNA from the or in the sample and hybridising and amplifying the DNA sequence using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 63; Column 97;
                                    Detection and quantification of specific fungi or bacteria useful e.g. in medical diagnosis and treatment of fungal and bacterial conditions, by hybridizing and amplifying DNA using sequences unique to the specie or group of species -
                                                                                                                                                                                                                                                                                                                                                                                            Fungus; bacteria; quantification; detection; infection; ulcer; asthma; allergy; sinusitis; probe; ss.
                                                                                                               WPI; 2002-462353/49.
                                                                                                                                           Haugland R,
                                                                                                                                                                                                                                              13-JUN-2000;
                                                                                                                                                                                                                                                                                                        US6387652-B1.
                                                                                                                                                                                                                                                                                                                                     Trichoderma viride.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                   15-APR-1998;
14-APR-1999;
                                                                                                                                                                                                                                                                           14-MAY-2002
                                                                                                                                                                                                                                                                                                                                                     Trichoderma
                                                                                                                                                                        (USSI ) US ENVIRONMENTAL PROTECTION AGENCY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                2000US-0593012
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                                                                                                                                         Vesper S;
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99US-0290990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.4; DB 24
Pred. No. 3.5e+03;
0; Mismatches 1
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Claim 64;

Column 97; 55pp;

English.

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ARESULT 13
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quantified in the samp
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                                                                                                                                                                                                                                                                                        Detection and quantification of specific fungi or bacteria useful e.g. in medical diagnosis and treatment of fungal and bacterial conditions, by hybridizing and amplifying DNA using sequences unique to the specie
                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Haugland
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Trichoderma atroviride.
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14-APR-1999;
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99US-0290990.
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Pred: No. 3.5e+03;
0; Mismatches 1;
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The invention relates to specific fungi or bacteria detected and quantified in a sample by extracting and recovering DNA from the organism in the sample and hybridising and amplifying the DNA sequence using sequences unique to the species/group of species but common to all isolates of the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe

Query Match Best Local Similarity

72.9**%**; 92.9**%**;

Score 12.4; Pred. No. 3.

4; DB 24; 3.5e+03;

Length

21;

With

Sequence 21

BP; 3 A; 7 C; 7 G; 4 T; 0 other;

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method

of the invention.

Claim 67;

Column 98; 55pp;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    groups or specified bacteria. The method enables detection and quantification of specific fungi or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment (fungi or bacteria associated with health problems such as infections, gastroenteritis, ulcers, asthma, allergies and sinusitis. It is also useful to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary haemorrhage from levels of Stachybotrys chartarum in air samples in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequences (ABN99429-ABN99653) for detection of specified figroups or specified bacteria. The method enables detection
                                       The invention relates to a method of detecting and quantifying fungi and bacteria; involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes
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                                                                                                                                                                                                                                             Detecting and quantifying fungi and bacteria, involves obtaining sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fungal detection; bacterial detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Trichoderma asperellum/hamatum probe.
                                                                                                                                                                                                  Claim 63; Page
                                                                                                                                                                                                                                                                                                                                         Haugland R,
                                                                                                                                                                                                                                                                                                                                                                                                     13-JUN-2000; 2000US-0593012.
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Trichoderma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          a asperellum.
a hamatum.
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                                                                                                                                                                                                  21; 110pp;
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                                                                                                                                                                                                    English.
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Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probe; primer;
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on and
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RESULT 15
ABK29219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Trichoderma asperellum. Trichoderma hamatum.
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                                                                                                                                                                                                                                                                                                                                                                                          Claim 64; Page 21; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-098078/13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fungal detection; bacterial detection; probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trichoderma asperellum/hamatum/viride
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK29219;
                                                                                                                                                                                                                           Sequence
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             23-APR-2002
                                      ABK29228;
                                                             ABK29228 standard;
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              (first entry)
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                                                             DNA;
                                                                                                                                                                                      72.9%;
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                                                                                                                                                                                                                           G; 4 T; .0 other;
                                                                                                                                                                          Score 12.4; DB 24; Pred. No. 3.5e+03; 0; Mismatches 1;
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                                                                                                                                                                                                  Length 21;
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 67; Page 22; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus; extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fungal detection; bacterial detection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 3 A; 7 C; 7 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-098078/13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             analysis
                                                                                                   Aspergillus ochraceus.
Aspergillus ostianus.
Aspergillus auricomus.
                                                                                                                                                                                                                       05-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (USSI ) US ENVIRONMENTAL PROTECTION AGENCY
                           13-JUN-2000; 2000US-0593012.
                                                                                                                                                     Fungus; bacteria; quantification; detection; infection; gastroenteritis;
ulcer; asthma; allergy; sinusitis; probe; 88.
                                                                                                                                                                                            Fungi probe SEQ ID NO 31.
                                                                                                                                                                                                                                                 ABN99459;
                                                                                                                                                                                                                                                                         ABN99459 standard;
                                                                           US6387652-B1
                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
13; Conserv
                                                                                                                                                                                                                                                                                                                                                               GTTGCTTCGGCGGG
                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         relates to a method of detecting and quantifying fungi and
                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                         72.9%;
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                                                                                                                                                                                                                                                                           ВР
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Pred. No. 3.5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                        24:
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15-APR-1998;

98US-081773P

14-APR-1999;

9905-0290990

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RESULT 18
ABK29056
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   quantified in a sample by extracting and recovering DNA from the organism of the sample and hybridising and amplifying the DNA sequence using sequences unique to the species/group of species but common to all solates of the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe sequences (ABN99429-ABN9963) for detection of specified fungal/fungal groups or specified bacteria. The method enables detection and quantification of specific fungal or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment of fungal or bacteria, sulcers, asthma, allergies and sinusitis. It is also useful to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary hamonary and some samples in Schener's asthma, allergies and sinusitis. It is also set to establish the risk of adverse health effects (e.g. pulmonary to be set to set t
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection and quantification of specific fungi or bacteria useful e. in medical diagnosis and treatment of fungal and bacterial condition by hybridizing and amplifying DNA using sequences unique to the spec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK29056 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                             Aspergillus
                                                                                                                                                                                                                                                                                                                                                                                                         Fungal detection;
Detecting and quantifying fungi and bacteria, in sequence of the fungus, extracting the DNA from
                                                                                                                                                                                            13-JUN-2000; 2000US-0593012
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                                                                                                                                                     ENVIRONMENTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ochraceus/ostianus/auricomus probe.
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                                                                                                       Vesper
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
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92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        specific fungi or bacteria detected and extracting and recovering DNA from the
                                                                                                                                                     PROTECTION
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                                                                                                                                                                                                                                                                                                                                                                                                         detection; probe; primer;
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Pred. No. 3.5e+03;
0; Mismatches 1
                                                                                                                                                     AGENCY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24;
                          involves obtaining
    the sample,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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S
                                    Matches
                                                  Best Local
                                                                 Query Match
                                                                                                                            The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                  Sequence 22 BP; 2 A; 8 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                   Claim 12; Page
                                                                                                                                                                                                                                                                                                                                                                    subjecting the DNA to
                                                                                                                                                                                                                                                                                                                                                    analysis
                                  13;
   -
                                                   Similarity
   GTTGCTTCGGCGGG 14
                                  Conservative
                                                                                                                                                                                                                                                                                                                   8; 110pp; English.
                                                  72.9%;
                                                                                                                                                                                                                                                                                                                                                                    polymerase
                                    0
                                  Score 12.4; D
Pred. No. 3.5e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                     chain
                                                                                                                                                                                                                                                                                                                                                                       reaction and
                                                   3.5e+03;
                                                                     8
                                                                 24;
                                      Indels
                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                       fluorescent
                                      0
                                                                                                                                                                                                                                                                                                                                                                     probe
                                    Gaps
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Q

GTTGCTTCGGCGAG

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RESULT 19
ABK29268
ID ABK29
XX
Fungal detection; bacterial detection; probe;
                                                                                                                                                                                         Aspergillus
                                                                                                                                                                                                    23-APR-2002
                                                                                                                                                                                                              ABK29268;
                                                                                                                                                                                                                        ABK29268 standard; DNA; 22 BP
                                                                                                                             13-JUN-2000; 2000US-0593012
                                                                                                                                       13-JUN-2001; 2001WO-US18892
                                                                                                                                                            WO200196612-A2
                                                                                                                                                                     Aspergillus ochraceus
                                                                                                                   (USSI ) US
                                                                                              2002-098078/13
                                                                                                         ,
                                                                                                                   ENVIRONMENTAL
                                                                                                                                                                                          ochraceus probe
                                                                                                                                                                                                    (first entry)
                                                                                                         ខ្ម
                                                                                                                   PROTECTION
                                                                                                                                                                                primer;
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The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes Detecting and quantifying fungi and bacteria, involves obtaining sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent F Claim 83; analysis Page 24, 110pp; English probe

in the

method of the invention.

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RESULT 20
ABK29271
IID ABK29271
IID ABK29271
AC ABK25
XX ABPE2
XX ABPE2
XX ABPE2
XX Pungs
XX 
RESULT 21
AAQ94376
ID AAQ94
XX
AC AAQ94
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Best Local S
Matches 13
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fungal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Haugland
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AAQ94376;
                                             AAQ94376 standard;
                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 84; Page 24; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-098078/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (USSI ) US ENVIRONMENTAL PROTECTION AGENCY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUN-2000; 2000US-0593012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUN-2001; 2001WO-US18892
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13; Conserv
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                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                             22
                                                                                                                                                                     GTIGCTICGCCGAG
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                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                           BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N
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                                                                                                                                                                                                                                                                                                                                                             A; 8 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bacterial detection; probe;
                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 72.9%;
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                                                                                                                                                                                                                                                                                  72.9%;
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                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                             G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.4; DB 24; Pred. No. 3.5e+03;
                                                                                                                                                                                                                                                                                    Score 12.4; DB 24
Pred. No. 3.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         σ
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                                                                                                                                                                                                                                                                                                        24;
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                                                                                                                                                                                                                                                                                                           Length
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                                                                                                                                                                                                                                                                                                             22;
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RESULT 22
AAV62524
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                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local,
                                                                                                                                                                                                                                                                                                                                                                                                           A novel method for the detection of plant pathogenic strains of fungi e.g. Septoria nodorum, S.tritici, Pseudocecrosporella herpotrichoides, Mycosphaerella fijiensis, M.musicola or Fusarium spp. involves the PCR amplification of sequences found in the Internal transcribed region (ITS) of the 185, 5.85 and 285 ribosomal. RNA genes by the primers AAQ94359-93 and AAT05357-72. These primers are desived from the ITS sequences of these fungi (AAT0534-T05404 and AAQ94398) and are strain specific. The amplification products of the reactions using these primers can be used with the capture primers AAT05378-93 in colourimetric assays. The primers and ITS DNAs can be used for the detection of specific fungal pathogen isolates and in monitoring diseas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Plant pathogen; fungus; Septoria nodorum; Septoria tritici; Pusari
Pseudocercosporella herpotrichoides; Mycosphaerella fijiensis; PCR
Mycosphaerella musicola; amplification; primer; ribosomal RNA gene
internal transcribed region; strain; capture; colourimetric assay;
                                     Internal transcribed spacer; ITS; ribosomal RNA; Fusarium avenaceum; Fusarium culmorum; Fusarium graminearum; Fusarium moniliforme; plant; Septoria avenae; Microdochicum nivale; Fusarium poae; fungal pathogen; PCR; nucleic acid detection; PCR primer: ss.
                                                                                                          M. fijiensis
                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-383005/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beck JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-APR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mycosphaerella
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                                                                                                                                                                                                                                                                                                                                                                                                   development in plant populations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page 15; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CIBA ) CIBA GEIGY AG
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                                                                                                                                   17-DEC-1998
                                                                                                                                                                                         AAV62524 standard;
                                                                                                                                                               AAV62524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding intervening transcribed sequence
                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fungal
                                                                                                                                                                                                                                                            GTTGCTTCGGGGGGCGAC
                                                                                                                                                                                                                                                                                       GTTGCTTCGGCGGGAAC
                                                                                                                                                                                                                                                                                                                                                                         20 BP; 1 A;
                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                          species
                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pathogens
                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                              71.8%;
                                                                                                                                                                                                                                                                                                                                                                         5 C;
                                                                                                          specific primer JB548.
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                                                                                                                                                                                                                                                                                                                               Score 12.2; DB 16
Pred. No. 4.5e+03;
                                                                                                                                                                                                                                                                                                                                                                          5
17
                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          other;
                                                                                                                                                                                                                                                                                                                                              DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used for detection
                                                                                                                                                                                                                                                                                                                                              Length
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                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                    disease
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Mycosphaerella fijiensis

Synthetic.

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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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19-APR-1995;
15-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequences AAV62507 to AAV62566 represent species specific PCR primers for various fungal isolates used for fungal detection in the course of the invention. The primers are designed based on the internal transcribed spacer (ITS) sequences of the various fungal species. The invention provides a DNA molecule isolated from the ribosomal RNA gene region of a fungal pathogen, where the DNA molecule consists of an ITS sequence selected from ITS1 and ITS2 of Fusarium culmorum, Fusarium graminearum, Fusarium moniliforme, Septoria avenae or Microdochicum nivale. A method for detecting F. graminearum, F. culmorum, F. moniliforme, F. poae, F. avenaceum and M. nivale isolates is also provided which comprises isolating DNA from a plant leaf infected with at least one of the above pathogens and amplifying parts of the ITS sequence of the pathogen(s) pCR using specific primers from within these sequences. The pathogen(s) are detected by visualising the amplified part
            20-APR-2000;
                                       20-APR-2000; 2000DE-1019967.
                                                                                               DE10019967-A1
                                                                                                                                                       protozoacide;
                                                                                                                                                                   Antibody; heavy chain; variable region; CDR3; human; gp96; cytostatic; complementarity determining region; antibacterial; virucide; fungicide
                                                                                                                                                                                                               Human gp96 antibody variable heavy chain CDR3 clone G12D DNA fragment.
                                                                                                                                                                                                                                             22-JAN-2002
                                                                                                                                                                                                                                                                          AAI68725
                                                                                                                                                                                                                                                                                                    AAI68725 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA isolated from fungal RNA, and its internal transcribed spacer sequence - used for detecting fungal pathogens in plant tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-541745/46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5814453-A
                                                                  25-OCT-2001
                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NOVS ) NOVARTIS FINANCE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the ITS sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                        GTTGCTTCGGCGGGAAC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 1 A; 5 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
            2000DE-1019967
                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-0887480.
95WO-US04712.
96US-0722187.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  71.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              56pp;
                                                                                                                                                                                                                                                                                                     略
                                                                                                                                                                                                                                                                                                                                                                              17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 12.2; DB 19;
Pred. No. 4.5e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                      fungicide;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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This invention describes a novel recombinant antibody (Ab) that binds continued and product of the invention has cytostatic, antibacterial, convincide, fungicide and protozoacide activity. Complexes of gp96 with the peptides of the invention induce a immune response (cytotoxic T cells) consectific for the associated peptides. Ab are used for purification or convince a immune response (cytotoxic T cells) converting of gp96, including intect complexes (C) with peptides (I), from converting amounts of tumour or infected cells. (C) are useful: (i) in human cytoxic and infections (by viruses, bacteria, mycoplasma, fungi and parasites (C) protozoa); and (ii) for raising anti-idiotypic antibodies. Ab make cytostable isolation of very pure gp96 in native form, even from small contaminate the final product by bleeding from the column). This sequence concamvalin A (as used in known processes and which may contaminate the final product by bleeding from the column). This sequence concamvalin action complementarity determining region (CDR3) fragment of the antibody raised against human gp96 described in two contaminates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Terness P,
Schild H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New recombinant antibody specific for native gp96, gp96 as complex with endogenous peptides, used for immunotherapy of e.g. tumor -
                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 17; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FENN-) FENNING BIOMED GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rammensee H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kleist C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Opelz G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Welschof M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arnold-Schild
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful for isolating autologous
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           ঠ
                          Matches
                                Query Match
Best Local
Φ
                          l Similarity
14; Conserv
            GTTGCTTCGGCGGGAAC
                          Conservative
                               71.8%;
22
             17
                          0
                                 Score 12.2;
Pred. No. 4.
                          Mismatches
                                4.5e+03;
                                       DB 24;
                                      Length
                          Indels
                                        24;
                          0
                          Gaps
                          0
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Sequence 24 BP; 3 A; 4 C; 8 G; 9 T; 0 other;

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RESULT 24
ABN99548
05-AUG-2002 (first entry)
                                                                                        13-JUN-2000; 2000US-0593012
                                                                                                                                             Penicillium brevicompactum.
Penicillium alberechii.
                                                                                                                                                                          Fungus; bacteria; quantification; detection; infection; gastroenteritis;
ulcer; asthma; allergy; sinusitis; PCR; primer; ss.
                                                                                                                                                                                                    Fungi PCR primer SEQ ID NO 120.
                                                                                                                                                                                                                                                             ABN99548 standard; DNA; 17
                         Haugland R,
                                                              15-APR-1998;
14-APR-1999;
                                                                                                           14-MAY-2002
                                                                                                                             US6387652-B1.
                                                                                                                                                                                                                                           ABN99548;
                                          (USSI ) US ENVIRONMENTAL PROTECTION AGENCY
                         Vesper S;
                                                              98US-081773P.
99US-0290990.
                                                                                                                                                                                                                                                               BP
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WPI; 2002-462353/49

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RESULT 25
ABK29145
ID ABK29
XX ABK29
XX ABK29
XX Penic
XX Funga
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences unique to the species/group of species but common to all isolates of the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe sequences (ABN99429-ABN9963) for detection of specified fungi/fungal groups or specified bacteria. The method enables detection and quantification of specific fungi or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment fungi or bacteria associated with health problems such as infections, gastroenteritis, ulcers, asthma, allergies and shus: its lis also useful to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary haemorrhage from levels of Stachybotrys chartarum in air samples in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection in medical by hybridi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 41; Column 94; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quantified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Penicllium brevicompactum. Penicllium alberechii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fungal detection; bacterial detection; probe; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Penicllium brevicompactum/alberechii forward PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quantified in a sample by extracting and recovering DNA from the order the sample and hybridising and amplifying the DNA sequence using
                                                                                                                                                      Detecting and quantifying fungi and bacteria, involve sequence of the fungus, extracting the DNA from the subjecting the DNA to polymerase chain reaction and
                                                                                                                                                                                                                                                                                                                     Haugland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK29145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK29145 standard; DNA; 17
                           The invention
                                                                              Claim 41; Page 15; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUN-2000; 2000US-0593012.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2001; 2001WO-US18892.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200196612-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               medical diagnosis and treatment of fungal and bacterial conditions, hybridizing and amplifying DNA using sequences unique to the species
                                                                                                                                                                                                                                                                      2002-098078/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12;
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                                                                                                                                                                                                                                                                                                                                                                             ENVIRONMENTAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP, 1 A;
                                                                                                                                                                                                                                                                                                                        Vesper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry
                           relates
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es to a method obtaining a se
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                                                                                                                                                                                                                                                                                                                                                                             PROTECTION AGENCY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         σ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         +
     sequence
     of detecting and quantifying equence of the microorganism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ngi or bacteria detected and and recovering DNA from the organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 24; I
                                                                                                                                                                                                                 involves obtaining a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17;
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                                                                                                                                                            sample, and fluorescent probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
        g fungi
to be
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RESULT 26
AAZ02872/
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Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABX29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                        PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY16734-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, epidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogramulomatosis. The polypeptides of the invention may be of use in treating these diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ02872 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ02872;
                                                                                                                                                                                                                                   Disclosure; Page 1560; 1755pp; English.
                                                                                                                                                                                                                                                                   Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                    WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                  17-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                             04-NOV-1998;
28-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9928475-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine; eye disease; conventional trachoma; nonendemic trachoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                98US-0107077.
97FR-0015041.
97FR-0016034.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           70.0%; Fi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       an ORF of Chlamydia trachomatis
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Pred. No. 5.7
O; Mismatches
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5.7e+03;
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RESULT 27
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                                                                                                  This invention describes a novel group of primer pairs for polymerase chain reaction (PCR) amplification of at least one genetic element (A), introduced by genetic modification of soya, maize or other plants, present in crude plant materials or derived compositions. The new primers and probes are used (i) to detect and/or quantify genetically modified plant materials in seeds, animal and human foodstuffs, especially where these have been processed so that DNA is extensively degraded and (ii) of for monitoring the amplification of (A). The probes may also be used to confirm a successful PCR amplification. The high annualing temperature of the primers ensures high specificity and sensitivity, even with badly degraded samples, and facilitates analysis by restriction analysis, since conly inexpensive restriction enzymes are required, reducing costs and comply inexpensive restriction enzymes are required, reducing costs and comply inexpensive restriction enzymes are required, reducing costs and comply inexpensive restriction enzymes are required, reducing costs and comply inexpensive restriction enzymes are required, reducing costs and comply inexpensive restriction enzymes are required, reducing costs and materials. AAR61845 represent the PCR primersused in the method of the invention which amplify the soybean (Glycine max) lectin gene, Zea mays invertase gene, Cauliflower Mosaic virus (caMV) 35s-promoter, and comply incomposin phosphotransferase) gene, Petunia hybrida comply incomposin phosphotransferase) gene, Petunia hybrida comply incomposin smodified Cryla(b) gene.
Query Match
Best Local S
Matches 13
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF61866 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New primer pairs for polymerase chain reaction, useful for detecting genetically modified plant materials in e.g. foods, have high annealing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-308749/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vollenhofer S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-OCT-2000; 2000WO-AT00283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200132919-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enol-pyruvyl-shikimate-3-phosphate synthase; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection; lectin; invertase; 35S-promoter; NOS terminator; NPTII; nopaline synthase; neomycin phosphotransferase; EPSPS; CrylA(b);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Soybean lectin PCR primer LECrl.
                                                                       Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            temperature for increased sensitivity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (OSTP ) OESTERR FORSCH SEIBERSDORF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amplification;
 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                         BP;
   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genetically modified organism; GMO; soya; maize; plant;
                                                                       3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mansfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
69.4%; Score 11.8; DB 22; 86.7%; Pred. No. 7.4e+03; tive 0; Mismatches 2;
                                                                       5 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 20; Length 20; 5.8e+03;
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                                  Length 19;
   Indels
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RESULT 29
ABI94529/c
ID ABI945
XX ABI945
AC ABI945
XX 16-FEB
XX Captur
XX KW Human;
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AAF69341/c
ID AAF693
XX AAF18e
XX AAF18e
XX AAF18e
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XX AAF18e
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to an antisense compound 8 to 30 bases in length targeted to the 5' untranslated (UTR) region, the coding region or the 3' UTR region human Integrin-linked kinase (hIK). The antisense oligonucleotides are useful for inhibiting the expression of human hIK in human cells or tissues, in vitro. The oligonucleotides can be utilized for diagnostics, therapeutics for the treatment of diseases associated with the expression of hIK, prophylaxis e.g. to prevent or delay infection, inflammation or tumor formation and as research reagent.
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        Human; K-ras;
                                                        Capture oligonucleptide Zip ID#1616 oligo #9.
                                                                                                                                                                     ABI94529;
                                                                                                                                                                                                                        ABI94529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antisense compounds capable of modulating expression of human Integrin-linked kinase, useful for diagnosis, prophylaxis and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-137069/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-APR-2001
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                                                                                                               16-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                     standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                     rectrerereenae 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 BP; 6 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Column 43, 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGCTTCGGCGGGAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e.g. tumors, associated with expression of the kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                            (first entry)
     PCR primer; probe; capture probe; mutation detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 11.8; DB 22;
Pred. No. 7.4e+03;
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AB185628/c
ID AB1856
XX AB1856
XX AB1856
XX TS-PBB
DT 15-PBB
DX Captur
XX CAPTUR
XX Human;
                                                                                                                                                                                                                                                                                                                                                                                                     C(I) have melting temperatures within a narrow range. The method is useful CC for detecting infectious diseases caused by bacterial infectious agents CC e.g. Salmonella, bisteria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and CC Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and polio virus, and parasitic infectious agents CC elected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and objectly defects. CC entecting cancer involving oncogenes, tumour suppressor genes, or genes CC involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC garticular papillomavirus types 16 and 18 and liver cancers. The CC method is also used for emvironmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning CC electron microscope and infrared microscope) the support at the cCC sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to CC fe the present invention.
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                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotide probes (I) for use on a support to which complem oligonucleotide probes (II) will hybridise with little mismatch, (I) have melting temperatures within a narrow range. The method:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-APR-2001; 2001WO-US10958
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                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-034366/04.
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                                       Capture oligonucleotide Zip ID#1616 oligo #1.
                                                                                                                                                       ABI85628 standard; DNA;
 Human; K-ras; PCR primer; probe; capture probe; mutation detection;
                                                                            15-PBB-2002
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                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                         TTGCTTCGGCGGGAA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zirvi M,
                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   BP; 7 A; 6 C; 5 G; 2 T; 0 other;
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                            invention.
                                                                                                                                                                                                                                                                                                                                            69.4%;
86.7%;
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                                                                                                                                                                                                                                                                                                                                              Score 11.8; DB 24; Pred. No. 7.4e+03;
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                                                                                                                                                                                                                                                                                                                              Mismatches
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CC (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents CC e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptcocccus neoformans, Candida albicans and CC Aspergillus fundgautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and pollo virus, and parasitic infectious agents CC medinesis. The method is also useful for detecting genetic diseases such CC electing cancer involving oncogenes, tumour suppressor genes, or genes CC involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, by 3 gene, human papillomavirus types 16 and 18 and liver cancers. The CC method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning CC electron microscope and infrared microscope) the support at the food sets occurred and correlating (using a computer) identified ligation to a gresence or absence of the target nucleotide sequences. AB182074 to CC of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-034366/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Barany F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-APR-2000; 2000US-197271P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-APR-2001; 2001WO-US10958
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             oligonuclectide probes (I) for use on a support to which complemoligonuclectide probes (II) will hybridise with little mismatch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 5; Fig 25; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CORR ) CORNELL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a method (M1) for designing
invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ementary
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Sequence 24 BP; 7 A; 7 C; 7 G; 3 T; 0 other;

present

á 밁 Matches Query Match Best Local 20 13; Ν Similarity TIGCTICGGCGGGAA 16 TIGCTICGGCIGCAA Conservative 69.4%; σ 0 Score 11.8; DB 2 Pred. No. 7.4e+03 Mismatches DB 24; <u>ب</u> Length 24; 0, Gaps

0

RESULT 31 ABI85629

ABI85629 standard; DNA; 24 ВÞ

ABI85629;

2X8X8X8X8 15-FEB-2002 (first entry)

Capture oligonucleotide Zip ID#1616 oligo #2.

Human; K-ras; PCR primer; probe; capture probe; mutation detection;

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RESULT 32
AAF80069/c
ID AAF800
XX
AC AAF800
XX
DT 11-JUN
XX
XX
DT PCR pX
XX
XX
DE PCR pX
XX
XX
DE PCR pZ
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC oligonuclectide probes (I) for use on a support to which complementary CC oligonuclectide probes (II) will hybridise with little mismatch, where CC (I) have melting temperatures within a narrow range. The method is useful CC for detecting infectious diseases caused by bacterial infectious agents CC e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus neoformans, Candida albicans and CC Aspergillus fundgautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and polio virus, and parasitic infectious agents CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. CC electing cancer involving oncogenes, tumour suppressor genes, or genes CC involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC grant feed industry, detecting comprises scanning (using e.g. a scanning CC and feed industry, detecting comprises scanning (using e.g. a scanning CC sets occurred and correlating (using a computer) intention to a presence or absence of the target nucleotide sequences. ABI92074 to CC fethe areaent invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            environmental monitoring; food
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-APR-2000; 2000US-197271P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200179548-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; Fig 25; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                             AAF80069 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CORR )
                                                                                                 11-JUN-2001
                                                                                                                                              AAF80069;
                                                 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002-034366/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24
                                                                                                                                                                                                                                                                                                                                                               TTGCTTCGGCGGGAA 16
                                                                                                                                                                                                                                                                                                                  TIGCTICGGCIGCAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zirvi M,
                                                 for a maize seed-specific gene promoter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP, 3
  seed-specific
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     69.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 C;
                                                                                                                                                                                             25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7
    promoter; maize;
                                                                                                                                                                                                                                                                                                                                                                                                       0.
                                                                                                                                                                                                                                                                                                                                                                                                            Score 11.8; DB:
Pred. No. 7.4e+0.
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 7 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Favis R, Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     .4e+03;
       fatty acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
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         metabolism;
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RESULT 33
ABQ61935/c
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is capable of driving transcription in a seed-preferred manner. The promoter is isolated from a gene which is homologous to the barley pZE40 gene. The seed-specific promoter is used for selectively expressing a nucleotide sequence encoding a polypeptide involved in fatty acid metabolism, protein metabolism, carbohydrate metabolism, phytohormone biosynthesis or in cell cycle regulation. The promoter may be used to produce transgenic plants. The genes of interest expressed by the promoter are also useful for varying the phenotype of seeds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein metabolism; carbohydrate metabolism;
cell cycle regulation; transgenic plant; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel seed-preferred promoter sequence useful for selectively expressing a first nucleotide sequence in a plant seed and fortransgenic plant production -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200121783-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zea mays
                                                                                                                                                                       Human, 88, PCR; primer; aquaporin; AQP5; AQP; water channel protein; oligonucleotide chip; OGN chip; cDNA chip; lung cancer; mutation detection; polymorphism detection; gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primers AAF80067-69 were used to amplify a maize promoter that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 19; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-257981/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jiao S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-SEP-2000; 2000WO-US26131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAR-2001
                                                                                                                                                                                                                            Human
                                                                                                                                                                                                                                                      03-OCT-2002
                                                                                                                                                                                                                                                                               ABQ61935;
                                                                                                                                                                                                                                                                                                       ABQ61935 standard;
                                                09-SEP-2000; 2000KR-0053821
                                                                       10-SEP-2001; 2001WO-KR01528
                                                                                                                         WO200220787-A1
                                                                                                                                                  Homo sapiens.
(GOOD-) GOODGENE INC (MOON/) MOON W.
                                                                                                 14-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                            aquaporin 5 (AQP5) gene oligonucleotide (OGN) chip PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Habben JE,
                                                                                                                                                                                                                                                                                                                                                                                              TTGCTTCGGCGGGAA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25 BP; 5 A; 8 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                       TTGCTGCGGCGCAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                    69.4%;
                                                                                                                                                                                                                                                                                                        25
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Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     .5e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            phytohormone biosynthesis; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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                                                                                                                                                                                                                              primer 674
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ARBSULT 34
AAP51334/C
ID AAP513
XX AAP513
XX AP513
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention comprises a mutant form of the human aquaporin 5 (AQP5) gene. Aquaporin (AQP) is a family of water channel proteins, through which water is transported into and out of cells - ten types of mammalian AQP have been identified so far. The invention also comprises an oligonucleotide (OGN) chip having 902 oligonucleotide primer sequences and a cDNA chip comprising one or more sequences from the human AQP5 gene is useful for disgnosing cancer (i.e lung cancer). The OGN chip is useful for detecting mutations and polymorphisms in AQP5 and the cDNA chip is useful for analysis of gene expression. The present DNA sequence represents a human aquaporin 5 (AQP5) gene oligonucleotide (OGN) chip PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Moon
Song
                                                                                                                                                                                                                                                                                                                    Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-393847/42.
                                                                                                                                                                                          21-JUN-2000;
                                                                                                                                                                                                                      28-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF51334;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF51334 standard;
               Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an ancleic acid that inhibits or reduces growth factor mediated
                                                                                                                                                            21-JUN-1999;
                                                                                                                                                                                                                                                 WO200078341-A1
                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                           neovascular condition of the
                                                                                                                                                                                                                                                                                                                                                                                                                                         IGP-I oligonucleotide #2294.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Z E
                                                                       2001-041421/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aquaporin 5 gene mutant useful for , prostate, or head or neck cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GTTGCTTCGGCGGGA 15
                                                                                                                                 MURDOCH
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Kim H,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                          2000WO-AU00693.
                                                                                                  Werther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                            99US-0140345
                                                                                                                                 CHILDRENS RES INST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA; 15
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86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 11.8;
Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .5e+03;
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RESULT 35
AAF51335/
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                                                                         Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an nucleic acid that inhibits or reduces growth factor mediate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Iraulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; lethyosis; serborrhoea; ruba; keratosis; neoplasia; seleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                      Amellorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisum nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation.
The present invention relates to a method for ameliorating the effects of skin disorders line method comprises contacting the skin with an antisense oligonucleotide, (for insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation,
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                                                                                                                                                                                                                             Example 8; Page 75; 201pp; English.
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                                                   A novel method for the detection of plant pathogenic strains of fungies. Septoria nodorum, S.tritici, Pseudocercosportella herpotrichoides, Mycosphaerella fijiensis, M.musicola or Fusarium spp, involves the PCR amplification of sequences found in the internal transcribed region (ITS) of the 185, 5.85 and 285 ribospomal RNA genes by the primers ARQ94359-93 and ANTO5357-72. These primers are derived from the ITS sequences of these fungi (ANTO5394-T05404 and ANQ94398) and are strain specific. The amplification products of the reactions using these primers can be used with the capture primers ANTO5378-93 in colourimetric assays. The primers and ITS DNAs can be used for the detection of specific fungal pathogen isolates and in monitoring diseas development in plant populations.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 15; 65pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                 in monitoring disease
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RESULT 38
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Matches 12
                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                        A novel method for the detection of plant pathogenic strains of fungite. 9. Septoria nodorum, S.tritici, Pseudocercosporella herpotrichoides, Mycosphaerella fijiensis, M.musicola or Fusarium spp, involves the PCR amplification of sequences found in the internal transcribed region (ITS) of the 185, 5.85 and 288 ribosomal RNA genes by the primers AAQ94359-93 and AAT05357-72. These primers are derived from the ITS sequences of these fungi (AAT05394-T05404 and AAQ94398) and are strain specific. The amplification products of the reactions using these primers can be used with the capture primers AAT05378-93 in colourimetric assays. The primers and ITS DNAs can be used for the detection of specific fungal pathogen isolates and in monitoring disease development in plant populations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Plant pathogen; fungus; Septoria nodorum; Septoria tritici; Fusarium; Pseudocercosporella harpotrichoides; Mycosphaerella fijiensis; PCR; Mycosphaerella musicola; amplification; primer; ribosomal RNA gene; internal transcribed region; strain; capture; colourimetric assay; isolate; development; population; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 15; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of plant fungal pathogens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA encoding intervening transcribed sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beck JJ, Ligon JM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-APR-1995;
                                                                                                                                                                                                                           Sequence 16 BP; 1 A; 4 C; 6 G; 5 T; 0 other;
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                                                                                                               12;
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                                                         GITGCTTCGGCGG
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                                                                                                                                    67.1%;
92.3%;
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Pred. No. 1.2e+04;
                                                                                                                                       Score 11.4; DB 16;
Pred. No. 1.2e+04;
                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      - used for detection
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RESULT 39 AAT55603/c

Stinchcomb DT, Chowrira B, Direnzo A, Draper KG, Dudycz L Grimm S, Karpeisky A, Kisich K, Matulic-Adamic J, McSwiggen JA, Modak A, Pavco P, Beigelman L, Sullivan SM; Sweedler D, Thompson JD, Tracz D, Usman N, Wincott FE; Woolf T;

Dudycz LW;

The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves relA mRNA at the nucleotide base position indicated in the DE line.

Ribozymes having modified bases and methods for use in inhibiting disease related genes

for

producing

them

Claim

2

Page

240; 407pp;

English.

WPI; 1995-351090/45.

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Intercellular adhesion molecule; rel A; tumour necrosis factor; INF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; philadelphia chromosome; inflammation; autoimmune disease; atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV; myocardial ischaemia; Kawasaki disease; septic shock; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human
AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enzymatic nucleic acid; ribozyme; trans cleavage; inh gene expression; downregulation; interleukin-5; IL-5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT55603 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                       23-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                         31-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09523225-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse relA hairpin ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-APR-1997
                                                                                                                                                                                                10-NOV-1994;
                                                                                                                                                    (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunodeficiency virus; acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                 94US-0316771.
94US-0319492.
94US-0321993.
                                                                                                                                                                                                                                                                                                                                               94US-0224483.
94US-0227958.
94US-0228041.
                                                                                                                                                                                                                                                                                                                             94US-0245736
94US-0271280
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94US-0292620
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          target sequence
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ARESULT 40
AAV62556
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local (
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                       Sequences AAV62507 to AAV62566 represent species specific PCR primers for various fungal isolates used for fungal detection in the course of the invention. The primers are designed based on the internal transcribed spacer (ITS) sequences of the various fungal species. The invention provides a DNA molecule isolated from the ribosomal RNA gene region of a fungal pathogen, where the DNA molecule consists of an ITS sequence selected from ITS1 and ITS2 of Fusarium culmorum, Fusarium
                                                                                                                                                                                                                                                                                                                                                                                                 DNA isolated from fungal RNA, and its internal transcribed spacer sequence - used for detecting fungal pathogens in plant tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JUL-1997;
19-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-SEP-1998
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graminearum,
                                                                                                                                                                                                                                                                                                                            Claim 3; Column 17; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-541745/46.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Internal transcribed spacer; ITS; ribosomal RNA; Fusarium avenaceum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fusarium species specific primer JB577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16 BP; 7 A; 5 C; 4 G; 0 U; 0 other;
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an VEGF-B exon an STAT3 phosph primer TEF-46. F2 antisense ol an TRAP100 PCR an FGF-BP gene-	Filmer #20 tor cys Mu expression vect junB gene antisens PCR primer used to	N epi	mer i	-40 n	prot	cumour speci scuPA deltal	M. tuberculosis as Zomomonas mobilis	o PCI	pric	agged	MN gene activity i	מסלא: אלאנייייייייייייייייייייייייייייייייייי	noty	nsec	haer	or D	rece	igon	rece	fic f	latp	ine /	2	berz	lAzyme	mmerl	1 17	1 17	1 17	117	117	plif;	hamme	merhe	rimer	Sequence of a muta
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14-MAY-2002
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                                                    15-APR-1998;
14-APR-1999;
                                                                                                       13-JUN-2000;
                                                                                                                                                                                                                                             ulcer;
                                                                                                                                                                                                                                                               Fungus; bacteria;
                                                                                                                                                                                                                                                                                               Fungi probe SEQ ID NO 188
                                                                                                                                                                                                                                                                                                                                                                      ABN99616;
                                                                                                                                                                                                                                                                                                                                                                                                      ABN99616 standard;
                   (USSI ) US ENVIRONMENTAL PROTECTION AGENCY.
                                                                                                                                                                                                                                     asthma; allergy; sinusitis;
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                                                                                                       2000US-0593012.
                                                    98US-081773P.
99US-0290990.
                                                                                                                                                                                                                                       quantification; detection; infection; gastroenteritis; lergy; sinusitis; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to specific fungi or bacteria detected and converting and sample by extracting and recovering DNA from the organism continues and in a sample by extracting and recovering DNA from the organism continues and probability and supplied the DNA sequence using continues unique to the species/group of species but common to all continues of the species/group of species. The method especially employs sequences of the species/group of species. The method especially employs continues and species and species and probe continues of specific fungal species and specific for specific on of specified fungal species and specific fungal species of continues of specific fungal species, useful medically for disgnosis and treatment of supplication of specific fungal medically for disgnosis and treatment of control to detect and/or quantify microorganisms in the environment e.g. constablish the risk of adverse health effects (e.g. pulmonary for bourses) as the samples in bourses health effects (e.g. pulmonary controls).
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Best Local
                            Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent proanalysis
                                                                                                                                                                                                                                                                                                                                                                                               Fungal detection; bacterial detection; probe; primer;
Claim 62; Page 20; 110pp; English
                                                                                                                                                                                                                         13-JUN-2000; 2000US-0593012
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                                                                                                                                                    Haugland R,
                                                                                                                                                                                                                                                         13-JUN-2001; 2001WO-US18892
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The invention relates to specific fungi or bacteria detected and quantified in a sample by extracting and recovering DNA from the organism in the sample and hybridising and amplifying the DNA sequence using sequences unique to the species/group of species. The method especially employs sequences selected from 225 fungal and bacterial primer and probe sequences (ANN99429-ABN99653) for detection of specified fungi/fungal groups or specified bacteria. The method enables detection and quantification of specific fungi or bacteria, or groups of fungal/bacterial species, useful medically for diagnosis and treatment of fungi or bacteria associated with health problems such as infections, quastroenteritis, ulcers, asthma, allergies and sinusitis. It is also
                                                                                                                                                                                                        Claim 36; Column 94; 55pp; English
                                                                                                                                                                                                                                       Detection and quantification of specific fungi or bacteria useful e.g. in medical diagnosis and treatment of fungal and bacterial conditions, by hybridizing and amplifying DNA using sequences unique to the specier or group of species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation, ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes
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Best Local
                                                                                                                                                                               The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABX29026-ABX29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                                                             Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting and quantifying fungi and bacteria, involves obtaining sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent property.
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RESULT 5
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ARESULT 5
ID ARESULT 5
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BARRING ARESULT 30-1
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                                                                                                                                Query Match
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Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell
                                                                                                                                                                                                                                                                    Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 8; Page 78; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proliferation and/or inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MURD-) MURDOCH CHILDRENS RES INST.
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                                                                                                                                                                                                                                                                                                                                       vessels or any other hyperplasia.
                                                                                                                                   13;
N
                                                                5 GCCCGGATCCAGGC 18
                                                                                                                                                                   Similarity
                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                    BP; 3 A; 5 C; 5 G; 2 T; 0 other;
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                                                                                                                                                                68.9%;
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                                                                                                                                Score 12.4; D
Pred. No. 1.3e
0; Mismatches
                                                                                                                                                                   .3e+04;
                                                                                                                                                                                                       DB
                                                                                                                                                                                            Length
                                                                                                                                                                                         , 15;
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                                                                                                                                   Gaps
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RESULT 6
AAF51792
ID AAF5

AAF51792 standard; DNA; 15

ВP

RESULT 7
AAL42362/c
ID AAL423
XX
AC AAL423
XX
AC AAL423
XX
AC AAL423
XX
AC AAL423

AAL42362

standard;

DNA;

24

28-JUN-2002 AAL42362;

(first

entry)

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GCCCGGATCCAGGC Conservative

Query Match Best Local : Matches

l Similarity 13; Conserv

68.9%;

Score 12.4; DB Pred. No. 1.3e+0 Mismatches

.3e+04;

Length

15; 0

Gaps

0

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AntiBense therapy; antiproliferative; antiinflammatory; antipBoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1, pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperneovascular condition of the retina; ss.
                                                                                      The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contracting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor, IGF binding protein [IGFBP]-2 or IGFBB3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina brain or skin or work for a retormandiared malitonancies of the
                                                                                                                                                                                                                                                                                                                                                                                            Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAR-2001
                                retina, brain or skin, growth factor-mediated malignancies, esclerotic disease, kidney disease, hyperproliferation of the blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JUN-2000; 2000WO-AU00693
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Sequence 15 BP; 2
                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 78; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JUN-1999;
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A; 6 C; 5
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  G; 2 T; 0 other;
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RESULT 0
ABA9657
ID ABA9
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XX ABA9
XX IP-M
XX Huma
XX Huma
XX Huma
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                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                 oestrogen, but does respond to tamoxifen. The fusion proteins are used to induce selective proliferation (in vivo or in vitro), of cells the invention are expected to contain an exogenous gene. The fusion proteins of the invention are particularly useful in gene therapy where haematopoletic stem cells are being targeted. Selective proliferation of of cells through fusion proteins allows effective gene therapy even where gene transfer efficiency is low. Also, since the fusion proteins can be expressed in various types of blood cells, the range of cells that can be targeted is enlarged. The present DNA sequence represents a PCR primer used in an example of the invention to amplify the coding sequence of the hormone binding domain (HBD) of the mouse 4-hydroxytamoxifen-responsive
                            Human; thyroglobulin 11; recombinant production; gene therapy; thyroid associated disease; goitre; thyroidist; hypothyroidism; hyperthyroidism; struma nodosa; adenomatous goitre; reverse transcription-PCR; RT-PCR primer; ss.
                                                                                               Human thyroglobulin 11 RT-PCR primer,
                                                                                                                               19-MAR-2002
                                                                                                                                                                                      ABA96574 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New fusion protein, useful in gene therapy for selective expansion transfected cells, comprises mutant estrogen receptor and cytokine receptor fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4-hydroxytamoxifen-responsive mutant oestrogen receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-305122/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA2309324-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytokine receptor; tamoxifen; selective haematopoietic stem cell; hormone bindin
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                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to fusion proteins which comprises a mutant
                                                                                                                                                                                                                                                              14
                                                                                                                                                                                                                                                                                         w
                                                                                                                                                                                                                                                                                                                                                                                                             oestrogen receptor.
                                                                                                                                                                                                                                                                                                                      . Similarity
                         transcription-PCR;
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                                                                                                                                                                                                                                                                                       GCGCCCGGATCCAG 16
                                                                                                                                                                                                                                                              GTGCCCGGATCCAG 1
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                                                                                                                                                                                                                                                                                                                                                                                BP; 5
                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         stem cell; hormone binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fusion protein;
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                                                                                                                                                                                                                                                                                                                                  68.9%;
                         RT-PCR primer;
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                                                                                                                                                                                                                                                                                                                                  Score 12.4; DB
Pred. No. 1.3e+C
                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                 SEQ
                                                                                                                                                                                                                                                                                                                                  1.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      oestrogen receptor;
proliferation; gene
ng domain; HBD;
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                                                                                                 NO:4.
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RESULT 9
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Matches 13
                                                                                                                                                                                                                                                                                                                                 encoding it (ABA96572), and a method for the recombinant production of thyroglobulin 11. The protein has a molecular weight of 11 kD. The present invention additionally discloses an antagonist of thyroglobulin 11 for therapeutic use, and an antibody which specifically binds to thyroglobulin 11. Thyroglobulin 11, and nucleotides which encode it may be used for treating thyroid-associated diseases, such as goitre, thyroiditis, hypothyroidism, hyperthyroidism, and struma nodosa (adenomatous goitre). The protein may also be used to screen for modulators of its activity or for peptide fingerprinting identification. The polynucleotide can be used as a primer for nucleic acid amplification reactions or as a probe for hybridiastion reactions, or in producing gene chips or microarrays. Sequences hange 573-ABA96574 represent reverse transcription-PCR (RT-PCR) primers used in an exemplification of the invention to isolate human thyroglobulin 11 cDNA.
 06-SEP-1999
                                                                       AAX59160 standard; DNA; 21
                                    AAX59160;
                                                                                                                                                                                                                                                                                                       Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to human thyroglobulin 11 (AAM53042), nucleic encoding it (ABA96572), and a method for the recombinant production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 17; 38pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human thyroglobulin 11 polynucleotide and polypeptide, useful diagnosis and treatment of goiter, thyreolits, hypothyreosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
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                                                                                                                                                                                                                                                 n 68.9%;
Similarity . 92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SHANGHAI BIOWINDOW GENE
                                                                                                                                                                                             ecceearccaeec 18
                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                     BP; 3 A; 11 C; 5 G; 5 T; 0 other;
(first entry)
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                                                                                                                                                                                                                                                 Score 12.4; DB 24
Pred. No. 1.3e+04;
                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                 DB 24;
                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                               Length
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Gaps

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arthritis; diabetic retinopathy; cytostatic; antinflammatory; extraggiogenic; antidiabetic; ophthalmological: Arthritis; primer; 88.
     09-DEC-1998;
                                WO9929858-A1
                                                      Synthetic
                                                                                                           Human soluble neuropilin-1 5'
                                              domo sapiens.
      98WO-US26138
                                                                                                           PCR primer.
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RESULT 10
ABK941
XX
ABK941
XX
ABK941
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ABK941
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ABK941
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ABK941
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Endoth
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Matches 14
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09-DEC-1997;
12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This oligonucleotide was used as 5' primer in the RT-PCR amplification of cDNA (see AXX59153) coding for a truncated, soluble human neuropilin-1 polypeptide (see AAY06319). The cDNA was cloned from a PC3 library. Neuropilin-1 is a novel vascular endothelial growth factor (VEGF) receptor that is expressed in endothelial cells and motile metastatic cancer cells. Claimed soluble neuropilin-1 polypeptides are useful for treatment of diseases or disorders associated with VEGF, e.g. metastasis, inappropriate angiogenesis, chronic inflammation, diabetic retinopathy and arthritis. Solid tumors can also be treated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gagnon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker; transgenic animal; solid support; cardiovascular regulator; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Soluble human neuropilins and related polynucleotides
                    Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CHIL-) CHILDRENS MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                             31-AUG-2001;
                                                                                                                             WPI; 2002-435060/46
                                                                                                                                                                                                                                                                            19-SEP-2000; 2000EP-0120123
                                                                                                                                                                                                                                                                                                                                                                               28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                WO200224747-A2
                                                                                                                                                                                                                             (EPID-) EPIDAUROS BIOTECHNOLOGIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ž
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14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Klagsbrun M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                             2001WO-EP10087.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0099615.
97US-0069155.
97US-0069687.
                                                                                                                                                                              Hoffmeyer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   67.8%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EDNRB) SNP detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer #15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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The invention describes a polynucleotide (I) of the endothelin (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) CC (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) CC (II) is useful for producing cells capable of expressing (I) co or (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (I), (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and conclecular variant gene comprising (I) is useful for identifying and conclecular variant of a polypeptide of modulating the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholesterolaemia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or bost cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 59; 190pp; English
Sequence 21 BP; 5 A; 3 C; 9 G; 3 T; 1 other;
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RESULT 11
                                                                                                                   Query Match
Best Local S
Matches 14
                 ABK94174;
                                   ABK94174 standard;
27-AUG-2002
                                                                                 20 CTTCGCCAGNATCCACGC
                                                                                                                    14;
                                                                                                  _
                                                                                                                            Similarity
                                                                                                  CTGCGCCCGGATCCAGGC 18
                                                                                                                    Conservative
(first entry)
                                   DNA;
                                                                                                                            67.8%;
77.8%;
                                     21
                                     ВP
                                                                                 w
                                                                                                                   0
                                                                                                              Score 12.2; Pred. No. 1.6e+04; O; Mismatches 4
                                                                                                                                      24;
                                                                                                                     4.
                                                                                                                                     Length
                                                                                                                     Indels
                                                                                                                                      21,
                                                                                                                     0
                                                                                                                     Gaps
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шинк; signaling system; cardiovascular disease; coronary heart dise hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholssterolaemia; forensic marker; transgenic Endothelin; single helin; EDN; endothelin converting enzyme; ECE; endothelin receptor; signaling system; cardiovascular disease; coronary heart disease; nucleotide animal; solid support; cardiovascular regulator; SNP

Endothelin

receptor B (EDNRB) SNP detection PCR primer #16.

Synthetic

WO200224747-A2

28-MAR-2002

31-AUG-2001; 2001WO-EP10087.

19-SEP-2000; 2000EP-0120123

(EPID-) EPIDAUROS BIOTECHNOLOGIE

Brinkmann U, Hoffmeyer

WPI; 2002-435060/46

Novel polynucleotide of the endothelin/endothelin converting

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RESULT 12
AAX24822/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC (EDN)/endothelin converting enzyme (EED)/receptors of EDN and ECE (EDNR) (endothelin converting enzyme (EED)/receptors of EDN and ECE (EDNR) CC signaling system which is associated with a cardiovascular disease. (I), CC the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing (I) CC or (III) is useful for producing cells capable of expressing a molecular CC variant polypeptide which is associated with a cardiovascular disease. (CC or (III), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing (I) a molecular variant gene comprising (I) is useful for identifying and obtaining a pro-drug or drug capable of modulating the activity of a molecular variant of a polypeptide of the EDN/EDR/ECE signaling system (CC or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDRR/ECE signaling system or its gene product. The isolated proteins and (CC polynucleotides encoding them are useful for preparation of a polynucleotides encoding them are useful for preparation of a coronary heart disease, hypertension, atherosclerosis, or related to abnormal angiogenesis or fatty acid metabolism e.g. diabetes and familial CC phypercholesteroleemia. The gene or a polynucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for CC comprising bolynucleotides, genes, vectors, polypeptides, antibodies or forention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding pathway. XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                               Synthetic
                                                                                                                                   17-JUL-1998;
                                                                                                                                                                   28-JAN-1999
                                                                                                                                                                                                                                                                              neurotransmitter;
                                                                                                                                                                                                                                                                                                                                      Oligonucleotide ZC15035 used in zsig37 DNA amplification.
                                                                                                                                                                                                                                                                                                                                                                                                         AAX24822;
              WPI, 1999-132244/11.
                                                                                                     18-JUL-1997,
                                                                                                                                                                                                  WO9904000-A1
                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                         Adipocyte-specific protein; homologue; zsig37; human; farty acid metabolism; energy balance; nutrition; ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               enzyme/receptors of endothelin and endothelin converting signaling system associated with cardiovascular disease, treating the disease -
                                                                       ( OMYZ)
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX24822 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                             sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14;
                                                                   ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Page 59; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTGCGCCCGGATCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67.8%;
nilarity 77.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B₽;
                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                    97US-0053154
                                                                                                                                    98WO-US14864
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                                                                                                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                    24
                                                                                                                                                                                                                                                                                                                                                                                                                                    ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.2; DB
Pred. No. 1.6e+C
O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4;
                                                                                                                                                                                                                                                                                           antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Oligonucleotide
ZC15721 (see AA)
                                                                                                                                                     New isolated adipocyte complement related develop products for modulating energy by protecting endothelial cells from injury
                                                                                                                              Example 4; Page 118; 128pp; English.
                                                                                                                                               neurotransmitter-modulated
                                                                                                                                              applications
                                                                                                                                                              related polypeptides nergy balance in mamma
                                                                                                                                                       o
R
                                                                                                                                                       for antimicrobial
                                                                                               with pr
1 866 bp
281g37
                                                                                                                                                                      used
                                                                                                              primer
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                                                                                               DNA
                                                                                                                                                                      ç
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(see AAX24813). The FCR product was used in the creation of mammalian expression vectors. zsig37 (see also AAW98013) can be used to modulate energy balance in mammals or to protect endothelial cells from injury. It can be used to modulate cellular metabolic reactions including adipogenesis, gluconeogenesis, glycogenolysis, lipogenesis, glucose uptake, protein synthesis, thermogenesis and oxygen utilisation. zsig37 may also modulate acetylcholine and/or norepinephrine release, and may find use as a neurotransmitter or as modulator of neurotransmission, as an antimicrobial agent or to modulate acetylcholine. Oligonucleotide ZC15035 was used as a primer, together ZC15721 (see AAX24821), in the PCR amplification of an fragment of human adipocyte-specific protein homologue (see AAX24813). The PCR product was used in the creati mammalian expression vectors. zsig37 (see also AAW9801 nutrient uptake.

Sequence 24 BP; 3 A; 6 C; ø G; 6 T; 0 other,

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밁
              S
                                      Query Match
Best Local (
                               Matches
20
               Н
                               14;
                               h 67.8%;
Similarity 82.4%;
14; Conservative
               CTGCGCCCGGATCCAGG
CCGAGCCCGGATCCATG
               17
4
                               0
                                       Score 12.2; DB 20;
Pred. No. 1.6e+04;
                               Mismatches
                              Indels
                                              Length
                                               24;
                              0
                              Gaps
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0

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ARAGULT 13
ARAG4122/6
ID ARAG4
XX ARAG4
AC ARAG4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              blood flow; thrombogenic activity; complement activity; angioplasty; collagen-mediated platelet adhesion; vascular reconstruction; aneurysm; coronary artery bypass graft; endarterectomy; microvascular repair; anastomosis; trauma; stroke; ischemia; reperfusion; endarterectomy; intestinal strangulation; cardiopulmonary bypass ischemia; recesitation; myocardial infarction; post-trauma vasopasm; accidental vascular trauma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  zsig37; adipocyte complement related protein; blood flow; thrombogenic activity; complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA64122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      percutaneous transluminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer for the human zsig37 DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 angioplasty; wound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             haemostasis inhibitor;
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WO200048625-A2 Homo sapiens.

17-FEB-2000; 2000WO-US04161 24-AUG-2000

19-FEB-1999; 22-NOV-1999; (ZYMO) ZYMOGENETICS 99US-0253604 99US-0444794 INC

Sheppard ğ, Lasser £ Bishop

2000-565335/52

Use of a human adipocyte complement related protein for promoting blood flow in the vasculature, pacifying damaged collagenous tissues, pacifying the surface of a prostatic biomaterial, and mediating wound repair -

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AAS19807
ID AAS19
XX AS19
XX AS19
XX AS19
XX OB-Ma
XX Human
Cance
KW Cance
KW Cance
KW Cance
XX Homo
XX Homo
XX Homo
XX II-AI
XX 
                                                  Genotyping human Ran GTPase activating protein 1 gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of the
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Similarity

66.7%;

Score 12; Pred. No.

DB 14; 2e+04;

Length

21;

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due to, e.g. vascular reconstruction (angioplasty, coronary artery bypass graft, endarterectomy, microvascular repair or anastomosis of a consciular graft), trauma, stroke or aneurysm. The polypeptide is also cuseful for pacifying damaged collagenous tissues due to injury associated with ischemia and reperfusion (trauma injury ischemia, intestinal strangulation, or injury associated with pre- and consciulation, myocardial infarction, or post-trauma vasopasm. In particular, post-trauma vasopasm includes stroke, percutaneous crassiluminal angioplasty, endarterectomy, accidental vascular trauma or surgical-induced vascular trauma. The polypeptide may also be used for pacifying the surface of a prostatic biomaterial, where the surface of the prostatic biomaterial is coated with collagen or collagen crassilumination fibrin or fibronectin. Finally, the polypeptide is useful for mediating wound repair, where the polypeptide enhances the progression of wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               zsig37 polypeptide an adipocyte complement related protein, and is an inhibitor of haemostasis and immune functions. The zsig37 protein is useful for promoting blood flow in the vasculature, where the polypeptide reduces thrombogenic and complement activity by the inhibition of the complement pathway and inhibiting collagen-mediated platelet adhesion, activation or aggregation. The methods are useful for treating injuries
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping
                                                                                                                                                                                                                                                                                                                       WO200179240-A2
                                                                                                                                                                                                                                                                                                                                                                                                 allele-specific
                                                                                                                                                                                                                                                                                                                                                                                                                   chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; genotyping;
cancer; irregular cell cycle associated disorder; ASO; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ASO primer #65 to detect human RANGAP1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS19807 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primers AAA64121-22 were used to amplify zsig37 gene fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; Page 93; 102pp; English
                                                                                                                                                                                                         17-APR-2000; 2000US-198072P
                                                                                                                                                                                                                                             17-APR-2001; 2001WO-US12455.
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                           2002-075068/10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
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                                                                                                                                                                     GENAISSANCE PHARM. INC
                                                                                                                               Choi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTGCGCCCGGATCCAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCGAGCCCGGATCCATG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                             JΥ,
                                                                                                                                                                                                                                                                                                                                                                                                 oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 6 C; 9 G; 6 T; 0 other;
                                                                                                                             Koshy B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          67.8%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.2; DB
Pred. No. 1.6e+0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24;
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Query Match
Best Local
                                                     Sequence 21 BP; 3 A; 8 C; 7 G; 3 T; 0 other;
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유 ð

Matches

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Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 0 A; 6 C; 6
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nucleotide sequence can be used in the detection of a determinant in pathogenic bacteria.
                                          plasmid DNA from clone pARC25 was used as a template to amplify the apyrase gene, by PCR, using the primers in AAQ40879-80. The apyrase enzyme is associated with the virulence of all species of Shigella (e.g. S. flexneri, S. dysenteriae, S. sonnei, S. boydii and related entero-invasive E.coli (EIEC) strains. The apyrase
                                                                                                                                                                                                                  ATP di:phospho:hydrolase or apyrase DNA sequence and prods. used for detection, therapy and protection against Shigella entero-invasive E.coli strains
                                                                                                                                                                                                                                                                                                                                                                                                                                         26-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GB2261878-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shigella; entero-invasive E.coli; EIEC; virulence; S. flexneri; S. dysenteriae; S. sonnei; S. boydii;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ40880 standard; DNA; 21
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                                                                                                                                                                       Example; Page 11; 40pp; English
                                                                                                                                                                                                                                                                                                          WPI; 1993-177505/22.
                                                                                                                                                                                                                                                                                                                                                   Ramachandran V, Roy RK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-JUN-1993.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ40880
                                                                                                                                                                                                                                                                                                                                                                                              (ASTR') ASTRA AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention relates to novel single nucleotide polymorphisms
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reverse primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92GB-0024716.
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85.7%;
                                                                                                                                                                                                                                                                                                                                                   Sankaran K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                     Subrahmanyam
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                                                                 boydii)
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RESULT 17
AAQ61767/c
ID AAQ617
XX AAQ617
AC AAQ617
XX DT 21-OCT
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DE HEV St
XX
KW Hepati
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AAQ64130
ID AAQ64430
XX AAQ64
XX AAQ64
XX AAQ71
XX AAQ71
XX AAQ71
XX AAQ71
XX Synth
XX Synth
XX WO941
XX I9-NC
XX 19-NC
XX 19-NC
XX I9-NC
XX IP-NC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A DNA sequence (AAQ64128) encoding a virulence-associated apyrase (ATP-diphosphohydrolase) in Shigella spp. and enteroinvusive Escharichia coli was located in S. flexneri plasmid pARC25. The 20 N-terminal amino acids of apyrase are given in sequence AAR54083 (positions 1-20 of the full apyrase sequence, AAR54082). The gene encoding apyrase was closed by PCR amplification of pARC25 DNA using the primers given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New DNA encoding virulence associated ATP disphosphoshydrolase - useful as hybridisation probe for specific detection of virulent Shigella sp. and enterosinvasive E. coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Apyrase; ATP-diphosphohydrolase; virulence; Escherichia coli; Shigella; probe; hybridization; plasmid pARC25; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 8 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 26; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI, 1994-199970/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ramachandran V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-JUN-1994.
  Hepatitis E
                                                   HEV strain BUR-121 primer D72.
                                                                                                       21-OCT-1994
                                                                                                                                                           AAQ61767;
                                                                                                                                                                                                            AAQ61767 standard; cDNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-NOV-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  CCGGATCCAGGC 18
                                                                                                                                                                                                                                                                                                                                              CCGGATCCAGGC 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
virus; HEV; strain SAR-55; open reading frame; ORF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                    (first
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                                                                                                    entry)
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 21
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AAT27478/c

ID

AAT27478 standard; DNA; 25 BF
XX

AC

AAT27478;

XX

DT

27-NOV-1996 (first entry)

XX

DT

27-NOV-1996 (first entry)

XX

HEV strain Burma-121 derived

XX

Hepatitis E virus; HEV; SAR-5

KW

structural region; antigen; c

KW

immunisation; infection; pri

KW

polymerase chain reaction; PC

XX

PN

W09610580-A2.

XX

PD

11-APR-1996.

XX

PD

03-OCT-1994; 94US-0316765.

XX

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PR

03-OCT-1994; 94US-0316765.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequences given in AAQ45198-200 and AAQ61687-777 are primers which were used in the isolation and amplification of the genomic sequence of the hepatitis E virus (HEV) strain SAR-55. These primers were based on sequences derived from the SAR-55 strain and a strain from Burma (BUR-121). The amplified sequence contains three open reading frames (ORFS). The proteins encoded by this sequence can be used to attimulate the production of protective antibodies upon injection into a mammal that would serve to protect the mammal upon challenge with wild type HEV. The proteins can be used for detection and diagnosis of HEV infection. This cDNA was isolated from primates innoculated with stool suspensions obtained from hepatitis E patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25 BP; 4 A; 6 C; 10 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Purified hepatitis E strain SAR-55 virus - used to develop prods. for use in detection, diagnosis, vaccines and therapy of hepatitis E virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Emerson SU,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9406913-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibody; detection; diagnosis; primates; stool suspension; amplify;
polymerase chain reaction; primer; burma; strain BUR-121; ss.
                                                                                                                                                                                                                                       Hepatitis E virus; HEV; SAR-55 strain; enteric transmission;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 40; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (USSH ) US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1994-118462/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 CCGGATCCAGGC 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7 CCGGATCCAGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                         Burma-121 derived forward primer 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Purcell RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DEPT HEALTH
                                                        95WO-US13102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0947263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93WO-US08849
                                                                                                                                                                                                                      antigen; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                        primer; Burma-121;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tearev
                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                        88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SA;
                                                                                                                                                                                                                         antibody; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                           (ORF-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 25
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DEPT

HEALTH & HUMAN SERVICES

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RESULT 19
AAV71688/c
ID AAV71
XX AAV71
XX AAV71
XX AAV71
XX AAV71
XX Hepat
XX Hepat
XX Hepat
XX Synth
OS Hepat
XX WO984
XX WO984
XX WO984
XX I1-AF
XX WPI Emert
XX WPI FR WPI FR
XX WPI FR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transmitted non-A, non-B hepatitis in Pakistan. The protein encoded by the structural region of the virus (i.e. ORF-2), which is capable of forming HEV like particles, is useful for the detection of HEV antibodies (pref. 19G or 19M) in blood, plasma, sera, cerebrospinal fluid, tissue, urine or pleural fluid. The protein, and anti-HEV antibodies generated using the protein, can also be used in vaccines for immunising an animal against HEV infection. The protein is identified as a band of greater than 50 kD following SDS-PAGE of cell lysates of insect cells infected with a HEV ORF-2 contg. baculovirus, i.e. the claimed recombinant expression vectors pPIC9-1779, -1780 and -1781.
                                                  New hepatitis E virus DNA from Pakistani strain SAR-55 - used for, e.g. developing products for diagnosis of, and vaccination against hepatitis E virus infection
        Example 1;
                                                                                                                                                           WPI, 1998-568733/48
                                                                                                                                                                                                                                                                                                                 11-APR-1997;
                                                                                                                                                                                                                                                                                                                                                              09-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                    22-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W09846761-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis E virus; HEV; SAR-55; diagnostic agent; vaccine; passive immunisation; open reading frame; ORF; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HEV ORF proteins encoding DNA amplifying primer D 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-FBB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV71688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV71688 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hepatitis E virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a hepatitis E virus (HEV) strain Burma derived primer, used in the isolation of the HEV strain SAR-55 CDNA. The HEV strain SAR-55 was implicated in an enterically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 44; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated and purified hepatitis E virus strain SAR-55 DNA - encodes antigenic protein useful in diagnosis, prophylaxis and treatment of hepatitis E virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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Page 46; 204pp; English
                                                                                                                                                                                                                                                              DEPT HEALTH & HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 6 C;
                                                                                                                                                                                                        Purcell RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Purcell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                 97US-0840316.
                                                                                                                                                                                                                                                                                                                                                              98WO-US07418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             꾶
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                                                                                                                                                                                                      Robinson RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12;
Pred. No.
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                                                                                                                                                                                                                                                           SERVICES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 17;
2e+04;
                                                                                                                                                                                                      Tsarev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                        SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 20
ADG68697/c
ID AAG686
XX Pullul
XX Pullul
XX EP6050
XX EP6050
XX EP6050
XX ABG686
XX AB
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reading frame (ORF) proteins ORF-1, ORR-2 and ORF-3. A host organism transformed or transferted with a recombinant expression vector containing the SAR-55 nucleic acid can be used to produce the HEV proteins, especially ORF-2 protein. The recombinant HEV proteins can be used as diagnostic agents and as vaccines for use against HEV infection. The detection of antibodies specific for HEV can be used for the diagnosis of infection and diseases caused by HEV, and for monitoring the progression of such disease. Such methods are also useful for monitoring the efficacy of therapeutic agents during the course of treatment of HEV infection and disease in a mammal. The antibodies can be used for detection or for passive immunisation of mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-DEC-1992;
15-JUL-1993;
19-NOV-1993;
                                                                      A probe was constructed which corresponded to the N-terminal part of the alkaline protease (AP) gene of Bacillus licheniformis SE2. The probe (BAQ68697) localised the N-terminal part of the gene to a 5.5kb PstI fragment as part of a procedure for preparing a deletion plasmid for deleting the AP gene from B.licheniformis SE2.
        Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                              New pullulanase from Bacillus deramificans - for starch saccharification etc., with good stability over wide temp. ranges, also related DNA vectors, transformed cells etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAV71605 to AAV71698 represent primers used for PCR amplification of the hepatitis E virus (HEV) DNA SAR-55 encoding frame (ORF) proteins ORF-1, ORF-2 and ORF-3. A host of the control of the 
                                                                                                                                                                                                                                                                                                                                       Example 15; Page 20; 61pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-210291/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUL-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP605040-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pullulanase; recombinant production; pullulan; amylopectin; hydrolysis; alpha-1,6-glycosidic bond; starch saccharificat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacillus licheniformis alkaline protease N-terminal probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             alkaline protease, ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ68697 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25 BP; 4 A; 6 C; 10 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SOLV ) SOLVAY SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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        BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    alpha-1,6-glycosidic bond;
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93BE-0000744.
93BE-0001278.
        N
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A; 6 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    66.7%; bu
100.0%; Pr
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Pred. No.
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0

Query Match

65.6%;

Score 11.8;

멺 15,

Length

18;

밁

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RESULT 22
AAT41772
ID AAT41
XX
AC AAT41
XX
DT 24-JJ
DT 24-JJ
XX
KW CDC2:
                                                                                                                                                                                                       Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                 in transformed Bacillus licheniformis, and related etc., used for pre-treatment of wood pulp to reduce ozone consumption in subsequent bleaching
                                                                                                                                                                                                                                                                                                                                                                                                                                        Amory A,
Ledoux P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Xylanase; endo-1,4-beta-D-xylanase; wood pulp; pulping;
biobleaching; bleaching; Bacillus licheniformis;
probe; hybridization; alkaline protease; ss.
                                                                                                                                                                                                                                                                          The C- and N-terminal portions of the Bacillus licheniformis SE2 alkaline protease gene were cloned by hybridization to the probes given in AAQ90831-32, respectively.
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-039214/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GB2279955-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ80932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ80932 standard;
                                                                                                                                                                                                                                                      Sequence 18 BP; 2 A; 6 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-1993,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-AUG-1995
CDC27; mitotic destruction complex; cell cycle regulatory protein;
                       Human CDC27 gene 5'
                                                24-JAN-1997
                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                                                                                                                                                                        Purified xylanase from Bacillus pumilus PRL B12 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-JAN-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alkaline protease N-terminus probe
                                                                       AAT41772;
                                                                                             AAT41772 standard; DNA; 24 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SOLV ) SOLVAY & CIE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-1993;
                                                                                                                                                         16 CTGCGCCAGGAGCCA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           μ
                                                                                                                                                                                14
                                                                                                                                                                                                        13;
                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                            23; Page 55; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTGCGCCCGGATCCA 15
                                                                                                                                                                                CTGCGCCCGGATCCA 15
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                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                    De
                      PCR primer.
                                               entry)
                                                                                                                                                                                                                  65.6%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86.7%;
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                                                                                                                                                                                                       Score 11.8; DB 16
Pred. No: 2.5e+04;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 2.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detroz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                              DB 16;
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    esp. produced
    ed DNA, vectors,
    uce chlorine or

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                                                                                                                                                                                                                             Length.
                                                                                                                                                                                                        Indels
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                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 23
ABA98282/c
                                                                                                                                                                                                                                                                                                                                                                                                     8
 밁
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primers (AAT41772 and AAT41773) were used to amplify the coding sequence (AAT41774) for CDC27 (AAM00364) from a human cDNA library. The amplified cDNA was cloned into vector pRSET and used to transfect E. coli JM109 for the prodn. of poly(His)-tagged CDC27 fusion protein. CDC27 can be used in novel cell-free ubiquitin-conjugating systems for the detection of inhibitors of ubiquitin-mediated proteolysis of cell cycle regulatory proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting inhibitors of ubiquitin-mediated proteolysis of CCRPs used in the treatment of proliferative and/or differentiation diseases, and in modulation of apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCRP; inhibitor; apoptosis; cell differentiation; primer; PCR, polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 3 A; 7 C; 10 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-485790/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           King RW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 24; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9633286-A1
 WPI; 2002-171436/22.
                                                                                  01-JUN-2000; 2000US-0585174.
                                                                                                                                  06-DEC-2001.
                                                                                                                                                         WO200192539-A2.
                                                                                                                                                                                 Pseudomonas mendocina
                                                                                                                                                                                                                   PHBA; para-Hydrobenzoate; liquid crystal polymer;
                                                                                                                                                                                                                                           Primer 48 for sequencing of pcu.
                                                                                                                                                                                                                                                                   18-JUN-2002
                                                                                                                                                                                                                                                                                         ABA98282;
                                                                                                                                                                                                                                                                                                                  ABA98282 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HARD ) HARVARD COLLEGE
                         Ramos-Gonzales
                                   Ben-Bassat A,
                                                                                                          22-MAY-2001; 2001WO-US16574
                                                                                                                                                                                                       toluene monooxygenase;
                                                          DUPO ) DU PONT DE NEMOURS &
                                                                                                                                                                                                                                                                                                                                                                                                                           13;
                                                                                                                                                                                                                                                                                                                                                                                                     3 GCGCCCGGATCCAGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                         1 65.6%; Score 11.8; DB 17;
Similarity 86.7%; Pred. No. 2.4e+04;
                                                                                                                                                                                                                                                                                                                                                                             GCGCGCGGATCCATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kirschner MW,
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                 (first
                     Cattermole M, MI, Ramos JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95US-0425299
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                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                   entry)
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                                                                                                                                                                                                        pcu gene; p-cresol; PCR primer; 88
                                                          S
                       Gatenby AA,
Sariaelani
                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                   Gibson KJ;
                                                                                                                                                                                                                      LCP;
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24; <u>.</u>

Gaps

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RESULT 24
AAZ26752/c
ID AAZ26
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Best Local :
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AAZ26752
                                                                                                                                                                                                                                                                                                                                                                                        Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allel especific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bcne marrow; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transformed with the p-cresol utilizing (pcu) genes are useful for producing para-hydroxybenzoate (PHBA) which can be used for synthesizing liquid crystal polymers (LCP). The sequences given in ABA98235-ABA98331 represent primers for; pcu sequencing, cloning of the p. putida pchC gene, sequencing tmox, cloning pcu for insertion into pMC3, construction of plasmids ppCUR1 and 2, mapping the transcript pMC3, construction of pcbA and B genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pacterial toluene monooxygenase enzyme pathway. The nucleic acid fragment encoding fragments of the invention may be used to isolate genes encoding proteins from the same or other microbial species. Bacterial strains transformed with the p-cresol utilizing from the same or other microbial species.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid fragments encoding bacterial toluene monooxygenase enzyme pathway, useful for isolating genes encoding proteins from the same or other microbial species, and for producing para-hydroxybenzoa
                                                                                                                                                                                                                                       20-MAR-1997;
                                                                                                                                                                                                                                                                                                  24-SEP-1998.
                                                                                                                                                                                                                                                                                                                                 WO9841648-A2
                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human polymorphic region 941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19
                                                                                                     diagnosis,
                                                                                                                  Identifying
                                                                                                                                               WPI; 1998-521232/44
                                                                                                                                                                             Housman
                                                                                                                                                                                                                                                                     19-MAR-1998;
                                                      Disclosure; Figure 7, 605pp; English.
                                                                                                                                                                                                           (VARI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CTGCGCCCGGATCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1
                                                                                                                                                                           Ö
                                                                                  ying target genes for allele-specific drugs - used for is, prevention and treatment of, e.g. cancers, atherosclerotic dysplastic lesions, endometriosis or graft versus host diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAGCACCCGTATCAAGGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A;
                                                                                                                                                                           Ledley FD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                       97US-0041057
                                                                                                                                                                                                                                                                     98WO-US05419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             64.4%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 C; 7
                                                                                                                                                                             Stanton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ŏ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genes encoding proteins from the for producing para-hydroxybenzoate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                       disease
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This invention describes a novel method for identifying an inhibitor potentially useful for treatment of cancer, where the inhibitor is active on a gene vital for cell growth or viability, and where the growth or viability, and where the growth or viability.

gene

for blocking the

metastatic cascade

This sequence represents a 3' splice site from the human CAPL gene, and is targeted by the oligonucleotide of the invention. The oligonucleotide inhibits expression of mRNA encoding CAPL (a calcium-binding protein of the S100 family) and has a sequence complementary to a sequence of the CAPL mRNA. The oligonucleotide may be used for inhibiting expression of CAPL mRNA in a cell-free reaction mixture or in a cell in vitro or for treating metastatic cancer. The oligonucleotide provides a novel method

Disclosure; Column 6;

28pp;

English

treating

metastatic

cancer

expression

of CAPL mRNA

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RESULT 25
ANV41150/c
ID ANV411
XX AV411
XX AV411
XX CAPL
DT 02-OCT
XX CAPL
DT 02-OCT
XX CAPL
XX CAPL
XX CAPL
XX CAPL
XX Homo c
XX D1-FEI
XX HYBR-FEI
XX HIGO
PT CREAT
XX D1scl.
XX D1scl.
XX D1scl.
XX This
CC CAPL

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV41150 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAPL; inhibitor; human; calcium-binding protein; S100 metastatic cancer; 3' splice site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV41150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
                                                                                                                                                                                                                                                                                                                      Oligo:nucleotides inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1998
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17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NORA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HYBR-)
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NORWEGIAN RADIUM HOSPITAL RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   splice
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTGCGCCCGGATCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8P;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Engebraaten
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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95US-0391375.
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77.8%;
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Pred. No. 3e
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RESULT 26
AAX17717 6
ID AAX17717 6
XX AX17717;
XX 10-MAY-195
XX US5877308
XX Human; CAI
XX W Human; CAI
XX OS Synthetic.
XX US5877308-
XX US5877308-
XX US5877308-
XX I6-FEB-19;
PR 19-FEB-19;
PR 19-FEB-19;
PR 19-FEB-19;
PR 03-MAY-19;
XX (HYBR-) H;
PA (NORA-) N(
XX I6-FEB-19;
PR 19-FEB-19;
PR 19-F
RESULT 27
ABK94171/c
ID ABK941
XX
AC ABK941
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AC ABK941
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Best Local S
Matches 14
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19-FEB-1995;
03-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention provides synthetic oligonucleotides which are complementary to and inhibit expression of human CAPL mRNA that includes the 3' or 5' splice site or the translational start site. The oligonucleotides prevent translation and splicing of RNA and can be used as antisense or triplehelix forming oligonucleotides to treat metastatic cancer.
  27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New synthetic oligonucleotides complementary to human CAPL mRNA - useful for inhibiting expression of s100 calcium binding protein the treatment of metastatic cancer
                                              ABK94171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-189725/16.
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Von Hofe E;
                                                                                          ABK94171
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(NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; CAPL; RNA splicing; antisense; metastatic cancer;
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                                                                                                                                                                                                                                        CTGCGCCCGGATCCAGGC 18
                                                                                                                                                                                                          CTGCTCCCAGATCCTGAC 1
                                                                                       standard;
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                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                             BP; 5 A; 3 C; 10 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Column 37; 27pp; English
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(first entry)
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95US-0391375.
96US-0642407.
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                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                        64.4%;
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77.8%;
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                                                                                                                                                                                                                                                                                                                   Score 11.6; DB 20;
Pred. No. 3e+04;
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Pred. No. 3e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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RESULT 28 ABK94172 ID ABK94 XX

ABK94172

standard; DNA; 21

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20 .

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CTGCGCCCGGATCCAGGC 18 CTTCGCCAGTATCCACGC

Matches Query Match Best Local (

Similarity

64.48;

Score 11.6; DB Pred. No. 3e+04; Mismatches

3e+04;

24;

Length Indels

21, 0

Gaps

0

Conservative

0;

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CC the gene encoding EDN, ECE or EDNR (III) or a vector (III) expressing (I), cor (III) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. CC (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and complexity of a molecular variant gene comprising (I) is useful for identifying and cobtaining a pro-drug or drug capable of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and companing system or its gene product. The isolated proteins and companing system or its gene product. The isolated proteins and composition for treating a cardiovascular disease such as coronary heart disease, hypertension, atherosclerosis, or related to comparising system are useful as cardiovascular disease of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support comprising polynucleotides, genes, vectors, polypeptides, antibodies or identify single nucleotide polymorphisms in DNA encoding pathway.

CC cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention describes a polynucleotide (I) of the endothelin (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) signaling system which is associated with a cardiovascular disease. (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atherosclerosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolaemia; forensic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 6; Page 59; 190pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin receptor B (EDNRB) SNP detection PCR primer #13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating the disease
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  21 BP;
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6 A; 3 C; 9 G; 3 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transgenic animal; solid support; cardiovascular single nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful treating the disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK94172;
                                                                                                                                                                                             Sequence 21
                                                                                                                                                                                                                                            comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a polynucleotide (I) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (EPID-)
                                                                                                  14;
N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Page 59; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EPIDAUROS BIOTECHNOLOGIE
                                                                                                                            Similarity
                                              CTGCGCCCGGATCCAGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           receptor B (EDNRB)
                                                                                                                                                                                                BP; 3 A; 9 C; 3 G; 6 T; 0 other;
                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hoffmeyer
                                                                                                                       64.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S
  19
                                                                                                0,
                                                                                                                       Score 11.6; DB 24;
Pred. No. 3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP detection PCR primer
                                                                                                  Mismatches
                                                                                                  4
                                                                                                                                               Length
                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    endothelin
                                                                                                  0,
                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for
                                                                                                  0
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RESULT 29

문 á

20

CTTCGCCAGCATCCACGC

Matches Query Match

14;

Conservative

o ;

Mismatches

Indels

0

Gaps

0

CTGCGCCCGGATCCAGGC 18

Local Similarity

64.4%;

Score 11.6; DB 24; Pred. No. 3e+04;

Sequence

21

BP;

ហ P

3 C; 10 G; 3 T;

0 other;

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ABK94175/c
IID ABK94175/c
IID ABK941
XX
AC ABK941
AC ABK
                                                                                                                                                                                                   CC (EDN)/pendothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR) CC signaling system which is associated with a cardiovascular disease. (I), CC the gene encoding EDN, ECE or EDNR (II) or a vector (III) expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (C) (II) is useful for producing cells capable of expressing a molecular variant polypeptide which is associated with a cardiovascular disease. (C) (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing and conclecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or the gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product, or for identifying and obtaining an inhibitor of the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE signaling system or its gene product. The isolated proteins and CC polymucleotides encoding them are useful for preparation of a polymucleotide composition for treating a cardiovascular disease such as CC comparation and angiogenesis or fatty acid metabolism e.g. diabetes and familial hypercholasterolasmia. The gene or a polymucleotide fragment of the EDN/ECE/EDNR signaling system are useful as forensic markers, for creating a transgenic animal and in creation of a solid support
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK94175 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK94175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel polynucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful treating the disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Endothelin receptor B (EDNRB)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2000; 2000EP-0120123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200224747-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes a polynucleotide (I) of the endothelin (EDN)/endothelin converting enzyme (ECE)/receptors of EDN and I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypertension;
                                                                 comprising polynucleotides, genes, vectors, polypeptides, antibodies or host cells of the invention. This sequence represents a PCR primer used to identify single nucleotide polymorphisms in DNA encoding cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               signaling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ension; atheroscierosis; angiogenesis; fatty acid mees; familial hypercholesterolaemia; foresnsic marker; enic animal; solid support; cardiovascular regulator nucleotide polymorphism; PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EDN; endothelin converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001WO-EP10087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           59; 190pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hoffmeyer S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      system; cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cardiovascular regulator; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    enzyme; ECE; endothelin receptor;
disease; coronary heart disease;
nesis; fatty acid metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #17.
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ABK94176 standard; DNA; 21

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CC (EDN) endothelin converting enzyme (ECE)/receptors of EDN and ECE (EDNR)
CC signaling system which is associated with a cardiovascular disease. (I),
CC the gene encoding EDN, ECE or EDNR (II) capable of expressing (I)
CC or (II) is useful for producing cells capable of expressing a molecular
CC variant polypeptide which is associated with a cardiovascular disease.
CC (II), (III), the EDN, ECE or EDNR polypeptide, or a cell expressing
CC a molecular variant gene comprising (I) is useful for identifying and
CC obtaining a pro-drug or drug capable of modulating the activity of a
CC molécular variant of a polypeptide of the EDN/EDNR/ECE signaling system
CC or its gene product, or for identifying and obtaining an inhibitor of
CC the activity of a molecular variant of a polypeptide of the EDN/EDNR/ECE
CC signaling system or its gene product. The isolated proteins and
CC polynucleotides encoding them are useful for preparation of a
CC comparising the encoding them are useful for preparation of a
CC comparished angiogenesis or fatty acid metabolism e.g. diabetes and familial
CC hypercholesterolaemia. The gene or a polynucleotide fragment of the
CC comprising polynucleotides, gene, vectors, polypeptides, antibodies or
CC comprising polynucleotides, gene, recetors, polypeptides, antibodies or
CC comprising polynucleotides, gene, vectors, polypeptides, antibodies or
CC consting a transgenic animal and in creation of a solid support
CC conformation of the invention. This sequence represents a PCR primer used
CC cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.

CC cardiovascular regulator proteins of the EDN/ECE/EDNR signaling pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polymucleotide of the endothelin/endothelin converting enzyme/receptors of endothelin and endothelin converting enzyme signaling system associated with cardiovascular disease, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endothelin; EDN; endothelin converting enzyme; ECE; endothelin receptor; EDNR; signaling system; cardiovascular disease; coronary heart disease; hypertension; atheroscierosis; angiogenesis; fatty acid metabolism; diabetes; familial hypercholesterolemia; forensic marker; transgenic animal; solid support; cardiovascular regula:or; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin receptor B (EDNRB) SNP detection PCR primer #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-AUG-2001; 2001WO-EP10087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 59; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-435060/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brinkmann U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-SEP-2000; 2000EP-0120123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200224747-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a polynucleotide (I) of the endothelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating
                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPID-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EPIDAUROS BIOTECHNOLOGIE AG.
               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hoffmeyer S;
                                                                              w
                                                                              Ą
                 64.4%;
                                                                              10 C;
                                                                              3 G; 5 T; 0 other;
Score 11.6; DB 24; Pred. No. 3e+04; 0; Mismatches 4;
                                     Length
  Indelia
                                         21;
0
Gaps
0
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ARESULT 31
ARAV1141/c
ID ARV41141/c
ID ARV41141/c
ID ARV41141/c
ID ARV41141/c
ID ARV41141/c
ID ARV41141/c
ID CAPL
XX CAPL
XX CAPL
XX CAPL
XX Metaat
XX Metaa
RESULT 32
AAV41142/c
ID AAV411
XX
AC AAV411
XC DT 02-OCT
XX
DE CAPL 3
XW CAPL;
KW CAPL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAPL; inhibitor; human; calcium-binding protein; $100 family; therapy; metastatic cancer; 3' splice site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CAPL 3' splice site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV41141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV41141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a 3' splice site from the human CAPL gene, and is targeted by the oligonucleotide of the invention. The oligonucleotide inhibits expression of mRNA encoding CAPL (a calcium-binding protein of the S100 family) and has a sequence complementary to a sequence of the CAPL mRNA. The oligonucleotide may be used for inhibiting expression of CAPL mRNA in a cell-free reaction mixture or in a cell in vitro or for treating metastatic cancer. The oligonucleotide provides a novel method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo:nucleotides inhibiting expression of CAPL \mathfrak{m}\mathtt{RNA} treating metastatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-OCT-1998
  CAPL; inhibitor; human; calcium-binding protein; $100 family; therapy; metastatic cancer; 3' splice site; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 5 A; 3 C; 11 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating metastatic cancer. The olig for blocking the metastatic cascade.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Column 6; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-446090/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Agrawal S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-FEB-1996;
17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-AUG-1998.
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                                                                                  CAPL 3' splice site.
                                                                                                                                         02-OCT-1998
                                                                                                                                                                                                  AAV41142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HYBR-) HYBRIDON INC.
(NORA-) NORWEGIAN RADIUM HOSPITAL RES FOUND.
                                                                                                                                                                                                                                                   AAV41142 standard; RNA; 22
                                                                                                                                                                                                                                                                                                                                                                                            18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CIGCGCCCGGATCCAGGC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTTCGCCAGCATCCACGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTGCGCCCGGATCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                            CTGCTCCCAGATCCTGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Engebraaten O,
                                                                                                                                         (first entry)
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95US-0391375.
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77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 11.6; DB 19;
Pred. No. 3e+04;
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RESULT 33
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Best Local
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                                              16-FEB-1996;
19-FEB-1995;
03-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents a 3' splice site from the human CAPI gene, and is targeted by the oligonucleotide of the invention. The oligonucleotide inhibits expression of mRNA encoding CAPI (a calcium-binding protein of the S100 family) and has a sequence complementary to a sequence of the CAPI mRNA. The oligonucleotide may be used for inhibiting expression of CAPI mRNA in a cell-free reaction mixture or in a cell in vitro or for treating metastatic cancer. The oligonucleotide provides a novel method
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                                                                                                                                                                                                                                          AAX17697 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Agrawal
                                                                                                                                                Synthetic.
                                                                                                                                                                   Human; CAPL;
                                                                                                                                                                                                       10-MAY-1999
                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                 treating metastatic cancer. The olig
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17-FEB-1995;
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Engebraaten
                                                                                                                                                                   RNA splicing; antisense; metastatic cancer;
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                                             96US-0602036.
95US-0391375.
96US-0642407.
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95US-0391375.
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                    RADIUM HOSPITAL RES FOUND.
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  Fodstad O,
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Pred. No. 3e+04;
0; Mismatches 4;
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RESULT 34
AAX17698;
ID AAX17698;
XX AAX17698;
AC AAX17698
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Best Local S
Matches 14
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19-FEB-1995;
03-MAY-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New synthetic oligonucleotides complementary to buseful for inhibiting expression of s100 calcium the treatment of metastatic cancer
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                                                                                                                                                                                                                               New synthetic oligonucleotides complementary to human CAPL mRNA useful for inhibiting expression of s100 calcium binding protein the treatment of metastatic cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-1996;
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                                                                                                                                                                                    Claim 2; Columns 27; 27pp; English.
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NORWEGIAN RADIUM HOSPITAL RES
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Pred. No. 3e+04;
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The invention provides synthetic oligonucleotides which are complementary to and inhibit expression of human CAPL mENA that includes the 3' or 5' splice site or the translational start site. The oligonucleotides prevent translation and splicing of RNA and can be used as antisense or triplehelix forming oligonucleotides to treat metastatic cancer.

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RESULT 35
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                                                                                                                                                                                                                                                 Thirty PRO
The present invention describes an isolated antibody that binds to one of the human PRO proteins designated PRO212, PRO290, PRO341, PRO355, PRO619, PRO717, PRO809, PRO8010, PRO819, PRO1005, PRO1005, PRO1009, PRO1025, PRO10130, PRO10097, PRO1107, PRO1111, PRO1153, PRO1182, PRO1184, PRO1025, PRO1281, PRO23, PRO394, PRO1117, PRO11710, PRO2094, PRO2145 OR PRO2198. PRO antagonist Pens used to inhibit tumour cell growth. The PRO polypeptides and nucleotides are useful in the treatment, diagnosis and prevention of cancer. The antibodies and other anti-tumour compounds maybe used to treat various conditions, including those characterised by overexpression and/or activation of the amplified antibodies and other compounds include benign or malignant tumours (e.g., renal, liver, kidney, bladder, breast, gastric, ovarian, colorectal, prostate, pancreatic, lung, vulva, thyroid, hepatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            proliferation; tumourigenesis; identification; cancer; PCR primer; hybridisation; probe; cytostatic; nootropic; neuroprotective; antiinflammatory; immunosuppressive; immunostimulant; antiangiogenic; leukaemia, lymphoid malignancy; neuronal disorder; glial disorder; estrocytal disorder; hypothalamic disorder; glandular disorder; astrocytal disorder; epithalamic disorder; stromal disorder; plandular disorder; epithalial disorder; stromal disorder;
                                                                                                                                                                                                                                                                                               WPI; 2000-572270/53.
                                                                                                                                                                                                                                                                                                                         Ashkenazi AJ,
Watanabe CK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-JAN-2000;
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                                                                                                                                                                                                                    Example 23; Page 133; 286pp; English.
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23-JUN-1999;
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99WO-US12252.
99US-0141037.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis; neoplastic disease; neoplastic cell growth;
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99US-0145698.
                                                                                                                                                                                                                                                                                                                                        Baker KP,
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                                                                                                                                                                                                                                                                                                                                      Goddard A,
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                                                                                                                                                                                                                                                 PRO polypeptides, useful of cancer -
                                                                                                                                                                                                                                                                                                                                        Gurney AL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22;
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                                                                                                                                                                                                                                                                                                                                        Roy MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 36
AAH78624/c
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Matches 14
                            Query Match
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Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    carcinomas, sarcomas, glioblastomas, and various head and neck tumours), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, stromal and blastocoelic disorders, and inflammatory, angiogenic and immunologic disorders. AAC58242 to AAC58366 represent PCR primers and hybridalsation probes used in the isolation of the human PRO sequences. AAC58367 to AAC58396 and AAB24057 to AAB24089 represent human.
                                                                                                           PCR primers AAH78624-25 were used to amplify exon 3 of the human reduced folate carrier (RPC) gene. Detection of mutations associated with the RPC gene are edd to assess between responsiveness of a cancer patient to antifolate chemotherapy. The RPC gene is associated with folate metabolism. The method is useful for assessing the responsiveness of a cancer patient to antifolate chemotherapy.
                                                                                     Sequence 23
                                                                                                                                                                                                                                                    Assessing responsiveness of a cancer patient to antifolate-containing chemotherapy, comprising searching for mutation or mutations in a gene associated with folate metabolism or uptake in cells derived from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; reduced folate carrier gene; RFC gene; cancer;
antifolate chemotherapy; folate metabolism; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH78624 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22
                                                                                                                                                                                                             Example;
                                                                                                                                                                                                                                                                                                                                       Assaraf Y,
                                                                                                                                                                                                                                                                                                                                                                                              06-MAR-2000; 2000US-0519673.
                                                                                                                                                                                                                                                                                                                                                                                                                         06-MAR-2001; 2001WO-IL00212
                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200165994-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer RFC780(9) for amplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH78624;
                                                                                                                                                                                                                                                                                                                                                                    (TECR ) TECHNION RES & DEV FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polynucleotide and protein sequences given in the exemplification present invention.
                                                                                                                                                                                                                                                                                                              2001-589905/66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 CTGCGCCCGGATCCAGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                              Similarity
                                                                                                                                                                                                          Page 59; 122pp; English.
  crececcedarccaesc 18
                                                                                   BP, 4 A; 8 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2 A; 10 C; 5 G;
                                                                                                                                                                                                                                                                                                                                        Drori S;
                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      64.4%;
                                                         64.4%;
                                           77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 11.6;
Pred. No. 3e
                                              Pred.
                                                         Score 11.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                              Mismatches
                                              ö
                                            3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BG
                                                         DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human RFC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21;
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                               Indels
                                                         Length
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                                                         23;
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                              Gaps
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AAF51790

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                                                                                                                                                                                                                                                         CC The present invention relates to a method for ameliorating the effects CC of skin disorders. The method comprises contacting the skin with an CC antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3), which is capable of cinhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonucleotide which can be used to design the antisense CC oligonucleotide of the present invention (see AAP45151 and CC oligonucleotides of the present invention (see AAP45151 and CC oligonucleotides of the present invention (see AAP45151 and CC oligonucleotides, pityriasis, ruba, pilaris, serborrhoea, keloids, CC psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, CC eratiosis, neoplasias, scleroderma, warts, benign growths, cancers of the CC skin, a hyperneovascular condition such as a neovascular condition of the CC retina, brain or skin, growth factor-mediated malignancies, other CC sclerotic disease, kidney disease, hyperproliferation of the inside of CC blood vessels or any other hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 37
                                                                     Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amellorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; Inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; neovascular condition of the retina; ss.
                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8, Page 78; 201pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUN-2000; 2000WO-AU00693.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MURD-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IGF-I oligonucleotide #2750
                                                                     l Similarity
12; Conserv
   Ç
                                                                                                                                                                                                                15
GCCCGGATCCAGG 17
                                                                                                                                                                                                         BP; 3 A;
                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                  63.3%;
                                                                                                                                                                                                         5 C; 5 G; 2 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Edmondson SR;
                                                                 0; Mismatches
                                                                                                  Score 11.4;
Pred. No. 3.
                                                                                                      3.8e+04;
                                                                                                                                DB 22;
                                                                 Indels
                                                                                                                            Length
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RESULT 39
AAV24277/c
ID AAV242

AAV24277 standard;

DNA;

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AAV24277;

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                                                                                                                                                                                                                                                                            The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an CC antisense oligonuclectide, (for Insulin-like Growth Factor [IGF]-1 CC receptor, IGF binding protein [IGFBP]-2 or IGFBP3], which is capable of CC inhibiting or reducing growth factor mediated cell proliferation, CC inflammation and/or other disorders. The present sequence is an CC oligonuclectide which can be used to design the antisense CC oligonuclectides of the present invention (see AAF45151 and CC oligonuclectides of the present invention (see AAF45151 and CC oligonuclectides of the present invention (see AAF45151 and CC oligonuclectides of the present invention (see AAF45151 and CC oligonuclectides of the present invention (see AAF45151 and CC oligonuclectides of the method is useful for ameliorating the effects of CC oligonuclectides of the present invention of the solution in the present invention of the condition of the skin, a hyperneovascular condition such as a neovascular condition of the cretina, brain or skin, growth factor-mediated malignancies, other condition or skin, growth factor-mediated malignancies, other the condition of the conditi
                                                                                                 Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor 1 receptor; IGF-1; pityriasis; IGF binding protein; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba, keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antis nucleic acid that inhibits or reduces growth factor mediated cell
                                                                                                                                                                                                            Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         proliferation and/or inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-041421/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wraight CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUN-2000; 2000WO-AU00693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200078341-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                 vessels or any other hyperplasia.
                                               σ
                                                                                                                             Similarity
                                    CCCGGATCCAGGC 18
CCCGGATTCAGGC 13
                                                                                                                                                                                                         BP; 3 A; 6 C; 4 G; 2 T; 0 other;
                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Werther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0140345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    78; 201pp; English.
                                                                                                                             63.3%;
92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Edmondson
                                                                                            Score 11.4; DB 22
Pred. No. 3.8e+04;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INST.
                                                                                                                                                    DB 22;
                                                                                                                                                 Length
                                                                                                                                                      15;
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RESULT 40
AAXOO122/c
ID AAXOO
XX AAXOO
XX AAXOO
XX 14-AE
XX Humar
XX Humar
XX Humar
XX Humar
XX Synth
OS Synth
OS Homo
XX Homo
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                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                           'New antibodies have been developed which are specific for human parathormone related peptides (hPTRP). The antibodies comprise chimeric L and/or H chains, where the C region is of human and L region of mouse, origin. The present sequence represents a PCR primer used in an example of the present invention. Host cells, transformed with vectors containing DNA encoding antibodies of the invention, can be used to produce the antibodies. The antibodies may be used to treat hypercalcaemia, especially that due to a malignancy, e.g. cancers of pancreas, lung, throat, larynx, tongue, gum, ossophagus, stomach, liver, breast, kidney, bladder, womb or prostate or malignant lymphoma. They may also be used for treatment of hypophosphaemia such as that due to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric; antibody; human parathormone related peptide; hPTRP; mouse; L chain; H chain; hypercalcaemia; cancer; malignant lymphoma; CDR; hypophosphaemia, pathogen; vitamin D resistance; V region; C region; humanised; PCR primer 88.
                                                                                                                                                                     AAX00122
                                                                                                                                                                                             122/c
AAX00122 standard; DNA; 19 BP.
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 19 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 3; Page 107; 182pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New chimeric antibodies against human parathormone related peptide(s) - useful for, e.g. treatment of hypercalcaemia and other disorders caused by malignant neoplasm(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-230640/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-APR-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9813388-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-1998
                              Synthetic.
                                                         Human; parathyroid hormone related protein; PTHrP; cachexia; cancer; inhibitor; humanised; PCR primer; ss.
                                                                                                        Human antibody
                                                                                                                                        14-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                      pathogens or to vitamin D resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sato K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JUL-1997;
26-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric antibody against hPTRP human L chain PCR primer MBCTLVR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHUS ) CHUGAI SEIYAKU KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   eapiens.
                sapiens
                                                                                                                                                                                                                                                                               13
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                                                                                                                                                                                                                                                                                                            GCCCGGATCCAGG 17
                                                                                                                                                                                                                                                                                                                                         63.3%;
nilarity 92.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry
                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97JP-0214168.
96JP-0255196.
                                                                                                        PCR primer MBC1LVR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-JP03382
                                                                                                                                                                                                                                                                                                                                                                                                       A; 6 C; 6 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yabuta
                                                                                                                                                                                                                                                                                                                                          <u>.</u>
                                                                                                                                                                                                                                                                                                                                         Score 11.4; DB 19;
Pred. No. 3.8e+04;
D; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 z
                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                         19;
                                                                                                                                                                                                                                                                                                                                          0
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RXGXGXKKKKKKKKKXXXDDDDDDDDDXX
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                            S
                                                           Matches
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                                                                                                                                         parathyroid hormone related peptide (PMRP) to its receptor, as an active component. This substance may be an antagonist to the receptor, or an antibody (preferably monoclonal) or antibody fragment, recognising PMRP. The antibody is preferably humanised or chimeric. The present invention also describes a humanised antibody prepared by hybridoma 23-57-137-1 (FERM BP-5631). The composition is used for the treatment of cachexia arising in connection with diseases such as cancer, thereby improving the quality of life of the patient. The present sequence represents a PCR primer used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                 Inhibitors of binding of parathyroid hormone related peptide to its receptor - useful for, e.g. treatment of cachexia arising from cancer or other diseases
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AX402921
AX402921.1 GI:21387912
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Method of identifying and quantifying specific fungi and
Patent: WO 0196612-A 262 20-DEC-2001;
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Sequence 262 from Patent
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Method of identifying and quantifying specific Patent: WO 0196612-A 244 20-DEC-2001;

UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; Fennellia.
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Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 241 20-DEC-2001;
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Aspergillus unguis
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceas; mitosporic Trichocomaceae; Asperg
                                                                                                                                                                                                                                                                                                                                                                              AX402775
Sequence 259 from Patent
AX402775
                                                                                                                                                                                                                                         Haugland, R. and Vesper, S.

Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 259 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
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Sequence 228 1
AX402744
AX402744.1 G:
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Method of identifying and quantifying specific fungi and Patent: WO 0196612-A 228 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
Location/Qualifiers
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Aspergillus caespitosus
Eukaryota; Fungi; Ascomycota;
Eurotiales; Trichocomaceae; m
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Similarity 87.5%;
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      195
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/db_xref="taxon:40381"
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|db_xref="taxon:176165"
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Pred. No. 2.3e+04;
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ltosporic Trichocomaceae; Aspergillus.
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AX402941/c
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Unclassified.
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1 (bases 1 to 15)
Haugland,R. and Vesper,S.
Method of identifying and quantifying specific fungi and bacteria Patent: US 6387652-A 195 14-MAY-2002;
Location/Qualifiers
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Trichoderma harzianum
Eukaryota, Fungi, Ascomycota, Pezizomycotina, So.
Hypocreales, mitosporic Hypocreales, Trichoderma
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Sequence 195 from Patent
AX402711
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Method of identifying and quantifying specific
Patent: WO 0196612A 195 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (
Location/Qualifiers
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                                                    Penicillium expansum.
Penicillium expansum
Eukaryota; Fungl; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales, Trichocomaceae; mitosporic Trichocomaceae; Penicillium
                                                                                                                                        Sequence 425 from Patent AX402941
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AR210283.1
Haugland, R. and Vesper, S. Method of identifying and quantifying specific fungi Patent: WO 0196612-A 425 20-DEC-2001; UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
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/db_xref="taxon:5544"
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Sequence 191
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Haugland,R. and Vesper,S.
Method of identifying and quantifying
Patent: US 6387652-A 194 14-MAY-2002;
Location/Qualifiers
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13; Conserv
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Haugland, R. and Vesper, S.
Method of identifying and quantifying
Patent: US 6387652-A 191 14-MAY-2002;
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AR210282
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/db_xref="taxon:27334"
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RESULT 15
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Sequence 194
AX402710
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Sequence 203
AR210291
Trichoderma asperellum
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreales; mitosporic Hypocreales; Trichoderma.
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                                                                                                                                                                                                                                                                                                                                                                                      Trichoderma asperellum

Bukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreales; mitosporic Hypocreales; Trichoderma.
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                                                                 AX402710.1
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Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 191 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 191 from Patent AX402707 AX402707.1 GI:21387698
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Haugland, R. and Vesper, S.
Method of identifying and quantifying specific fungi and bacteria
Patent: US 6387652-A 203 14-MAY-2002;
Location/Qualifiers
                                        Trichoderma asperellum.
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7 c 7 g
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AR210119
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Sequence 31
AR210119
                                                    1 (bases 1 to 22)
Haugland,R. and Vesper,S.
Method of identifying and quantifying
Patent: US 6387652-A 31 14-MAY-2002;
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AX402719
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Method of identifying and quantifying specific fun
Patent: WO 0196612-A 203 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trichoderma viride.
Trichoderma viride
Eukaryota; Fungi, Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreales; mitosporic Hypocreales; Trichoderma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Haugland,R. and Vesper,S.
Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 194 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
Location/Qualifiers
                                                                                                      Unclassified.
                                                                                                                 Unknown
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Similarity 92.9%;
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        organism="unknown"
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Pred. No. 4e+04;
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Sequence
AX402547
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Detection of fungal pathogens using the Patent: US 5814453-A 24 29-SEP-1998;
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Aspergillus ochraceus
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Asperg
                                                                                                                                                                                                                                                                                  AR043140
Sequence 24 from patent US
AR043140
AR043140.1 GI:5964148
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Method of identifying and quantifying specific
Patent: Wo 0196612-A 31 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (
Location/Qualifiers
                                                                                                                                                                                                                                                 Unknown
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                                                     GTTGCTTCGGGGGCGAC
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/db_xref="taxon:40380"
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Pred. No. 4e+04;
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Pred. No. 5.2e+04;
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Ligon, J.M. and Beck, J.J.

Ligon, J.M. and Beck, J.J.

Detection of fungal pathogens using the polymerase chain

Detection of fungal pathogens, 1999;

Patent: US 5955274-A 24 21-SEP-1999;

Location/Qualifiers
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Sequence :
132080
132080.1
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            Terness, P., Kleist, C., Opelz, G., Welschof, M. Schild, H. and Rammensee, H.G.
Antibodies against native gp96, production Patent: WO 0181423-A 15 01-NOV-2001;
Dr. Fenning Biomed GmbH (DE)
                                                                                                                                                             AX286541
Sequence 15 from Patent
AX286541
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Ligon, J.M. and Beck, J.J.
Detection of fungal pathogens using the polymerase chain reaction
Patent: US 5585238-A 24 17-DEC-1996;
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                                                                                                        synthetic construct.
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Location/Qualifiers
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                                                                 Welschof, M.,
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   ch 70.6%; Score 12; DB 1 Similarity 100.0%; Pred. No. 6.0 12; Conservative 0; Mismatches
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Sequence
AR210208
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Penicillium brevicompactum
Penicillium brevicompactum
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae; Penicillium.
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                                                                                                               Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 120 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
Location/Qualifiers
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12; Conserv
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                                                                /organism="Penicillium
/db_xref="taxon:5074"
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/db_xref="taxon:32630"
/note="Zufallssequenz, codiert fur CDR3 einer schweren
Kette eines Antikorpers, Herstellung siehe Nissim et al,
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AR125569/c
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18
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Sequence 70
AR125569
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Bennett, C.Frank. and Cowsert, L.M.

Antisense modulation of integrin-linked kinase

Patent: US 6177273-A 70 23-JAN-2001;

Location/Qualifiers
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                                               l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Glycine max
Eukaryota; Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papillonoideae; Phaseoleae;
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/organism="Glycine max"
/ob_xref="taxon:3847"
/note="Sojalektin-Gen LE1"
5 g 6 1
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Pred. No. 8.9e+04;
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/db xref="taxon:32630"
/note="Hypothetical Probe Sequence"
/ c 7 c 7 g 3 t
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/db_xref="taxon:32630"
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from Patent WO0121783
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Pred. No. 8.9e+04;
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AR043180
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AR043173
AR043173.1 (
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Sequence 64
AR043180
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1 (bases 1 to 25)

Jiao, S., Habben, J.E. and Niu, X.
Seed-preferred promoter from maize: Seed-preferred promoter from maize: Patent: WO 0121783.4 4 29-MAR-2001; PIONEER HI-BRED INTERNATIONAL, INC. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 16)
Beck, J.Joseph.
Detection of fungal pathogens using the polymerase chain reaction
Patent: US 5814453-A 57 29-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13,
                                                                                                                                                                                                                                                                                               12;
                                                        Beck, J. Joseph.

Detection of fungal pathogens using the Patent: US 5814453-A 64 29-SEP-1998;
                                                                                                                  Unknown
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/db_xref="taxon:4577"
8 c 6 g
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4 c 6 g
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Sequence 64
AR074680
AR074680.1
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12; Conserv
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Ligon, J.M. and Beck, J.J.
Detection of fungal pathogens using the polymerase chain reaction
Patent: US 5955274 A 64 21-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
AR074673
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Ligon, J.M. and Beck, J.J.
Detection of fungal pathogens using the
Patent: US 5955274-A 57 21-SEP-1999;
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12; Conserv
                                                                                                                 Similarity 92.:
12; Conservative
                                                                                                                                                                                                                                             Unclassified.
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Pred. No. 1.5e+05;
0; Mismatches ]
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Pred. No. 1.5e
0; Mismatches
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S 5955274.
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I62258/c
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Query Match
Best Local Similarity
Matches 12; Conserv
                                                          1 (bases 1 to 16)
Stinchcomb,D.T., Draper,K.G. and McSwiggen,J.
Rel a targeted ribozymes
Patent: US 5658780-A 812 19-AUG-1997;
                                                                                                                                                                                                     Sequence 812 from patent I62258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Haugland, R. and Vesper, S.
Method of identifying and quantifying s
Patent: WO 0196612-A 198 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION
LOCATION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Trichoderma longibrachiatum
Trichoderma longibrachiatum
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Hypocreales; mitosporic Hypocreales; Trichoderma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 198 from Patent AX402714
                                                                                                                         Unclassified.
                                                                                                                                                        Unknown.
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Haugland, R. and Vesper, S.
Method of identifying and quantifying
Patent: US 6387652-A 198 14-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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nilarity 92.3%;
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organism="unknown"
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/db_xref="taxon:5548"
5 c 6 g
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1 (bases 1 to 18)
Kikuchi, Y., Kiyokawa, S., Shimada, Y.,
Okinaka, Y.
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Sequence 19 from patent
AR151759
AR151759.1 GI:15117809
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Location/Qualifiers
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Kikuchi,Y., Kiyokawa,S., Shimada,Y., Ohbayashi,M.,
Okinaka,Y.
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Goshorn, S.C., Graves, S.S., Schultz, J.E., Lin, Y., Sanderson, J.A.
                                                                                                                                                                                 Streptavidin expressed gene fusions and methods of Patent: WO 0075333-A 32 14-DEC-2000; NEORX CORPORATION (US)
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Sequence 32 from Patent WO0075333.
AX057972
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Hamelin,R.C.
Detection of plant pathogenic fungi
Patent: US 5792611-A 1 11-AUG-1998;
Location/Qualifiers
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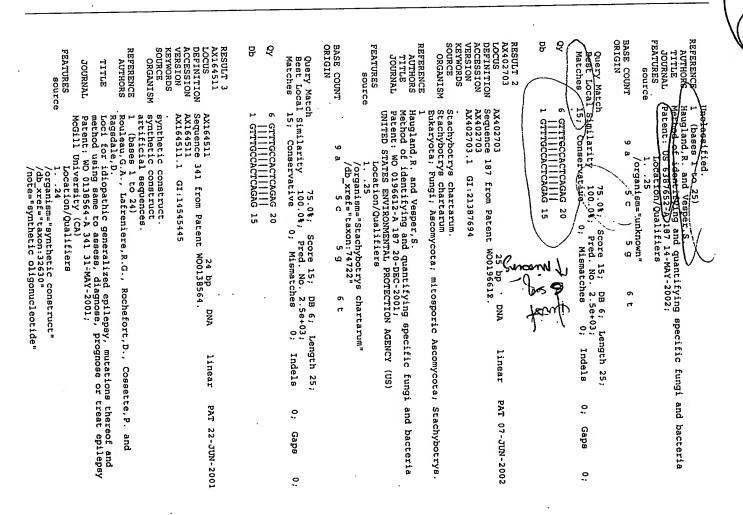
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1 (bases 1 to 17)

Draper,K.G., Dadyktz,L.W., M.
Holesek,J.J. and Mamone,A.J.
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Ward, D.T. and Cowsert, L.M.
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Patent: US 6165728-A 17 26-DEC-2000;
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07/882823,14-MAY-1992 US
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07/88289,14-MAY-1992 US
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KENNETH G DRAPER, LEC W D
WAYSEJAK,
PI JAMES J HOLESEK, ANT
PC C12N15/09, C12N5/10,
PC A61K39/135,
PC A61K39/145, A61K39/2
PC A61P31/14, A61P31/16
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PC C12N15
C12N15/00,
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Method and reagent for inhibiting viral replication Patent: JP 2000342286-A 118 12-DEC-2000;
RIBOZYME PHARMACEUTICALS INC
OS Artificial Sequence
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07-DEC-1992 US 07/987130,07-DEC-1992 US 07/
KENNETH G DRAPER,LEC W DADYKTZ,JAMES A MACSWIGEN,
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C12R1:93)
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C12N15/09, C12N5/10, C12N7/00, C12N9/22//(C12N5/10, C12R1:91),
                                                                                                JAMES J HOLESEK,ANTHONY J MAMONE
C12N15/09,C12N5/10,C12N7/00//A61K38/43,A61K39/125,A61K39/13,
A61K39/135,
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2N15/09,C12R1:93),C12N15/00,C12N5/00,A61K37/48,(C12N15/00, I
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07/88436, 14-MAY-1992 U
07/9833128, 26-AUG-1992 U
07/9633128, 07-DEC-1992 U
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Sequence 3 from Patent W00140471.
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                                                                                                                              Mycoplasma mycoides.
Mycoplasma mycoides
Bacteria, Firmicutes; Mollicutes; Entomoplasmatales;
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                                                        Frey, J.J., Nicolet, J.J. and Abdo, E.E. Antigenic protein lppq of Mycoplasma sc., its preparation and use patent: WO 0140471-A 3 07-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           synthetic construct.
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/organism="Mycoplasma mycoides"
/db_xref="taxon:2102"
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Pred. No. 4.1e+04;
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AX353540.1 GI:18618615
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/db xref="taxon:32630"
/note="lower primer FVR2536"
/note="5 c 5 g 4 t
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Sequence 393 from Patent
AX422057
AX422057.1 GI:21525439
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Patent: WO 0188124-A 393 22-NOV-2001;
RIBOZYME PHARMACEUTICALS, INC. (US) ; GLAXO (
                                                                                                                                                                                                                                                                                                             Homo sapiens
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Patent: WO 0159103-A 2611 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  synthetic construct
artificial sequences.
1 (bases 1 to 17)
Blatt, L., Mcswiggen, J. and Chowrira, B.
Method and reagent for the modulation
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/db_xref="taxon:32630"
/note="Nucleic Acid"
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1124 Columbia; Mailstop M318
Seattle, WA 98104; USA
e-mail: EAOStrander@lbl.gov
PCR Buffer: PCR buffer (Perkin-Elmer/Cetus)
PCR Profile: Denaturation: 94 degrees C for 1.00
Annealing: 55 or 59 degrees C for 0.45 minutes
Polymerization: 74 degrees C for 1.00 minutes
                                                                                         1 (bases 1 to 22)
Goldgaber, D.Y., Schwarzman, A.L. and E
Methods for the detection of soluble
(.beta.AP) or soluble transthyretin (
Patent: US 5744368-A 2 28-APR-1998;
                                                                                                                                                                                                                             Sequence 2
AR003705
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Ostrander, E.A., Mapa, P.A.,
One hundred and one new si
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Mammalia; Eutheria;
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PCR identification;
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                                                                                                                                                           Unclassified
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/tlssue_type="spleen"
/desue_fadult"
/tlssue_lib="E. Ostrander, i.
complement(1. .20)
a 5 c 5 g 6 t
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                                     organism="unknown"
2 c 10 g
                                                                           ocation/Qualifiers
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61.0%;
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Pred. No. 8.6e+04;
0; Mismatches 3;
 Score 12.
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AX402624
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AR040151
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14; Conserv
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Haugland, R. and Vesper, S.
Method of identifying and quantifying
Patent: US 6387652-A 108 14-MAY-2002;
Location/Qualifiers
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Sequence 108 from Patent
AX402624
AX402624.1 GI:21387615
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                                                                                                                                                                                              Haugland,R. and Vesper,S.
Method of identifying and quantifying specific
Patent: WO 0196612-A 108 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (
Location/Qualifiers
                                                                                                                                                                                                                                                                        nematode egg-parasitic fungus.
Paecilomyces lilacinus
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Eurotiomycetes;
Eurotiales; Trichocomaceae; mitosporic Trichocomaceae;
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nilarity 82.4%;
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4 c 6 g
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/db_xref="taxon:33203"
4 c 6 g 5 t
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Pred. No. 8.7e+04;
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Pred. No. 8.7e+04;
D; Mismatches 3
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AR166679
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Stinchcomb,D.T. and McSwiggen,J.A.
Interleukin-2 receptor gamma-chain ril
Patent: US 5807743-A 999 15-SEP-1998;
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AR166679
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Hess,J.W., Caskey,C.Thomas, Liu,Q. and Phillips,M.Sean.
Rat ob-receptors and nucleotides encoding them
Patent: US 6281346-A 29 28-AUG-2001;
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Sequence 22 from Patent
AX268543
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                                                         Welcher, A.A. and Calzone, F.J.
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                                      Cd20/ige-receptor like molecules and uses thereof patent: WO 0174903-A 22 11-OCT-2001;
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3 c 7 g
/organism="synthetic
                   Location/Qualifiers
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Pred. No. 1.1e+05;
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AX059295/c
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Blatt, L., Mcswiggen, J. and Chowrira, B.M.

Method and reagent for the modulation and dia
nogo gene expression
Patent: WO 0159103-A 2000 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M. McSwiggen, James (US);
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Sequence 2000 from Patent
AX216558
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synthetic construct
artificial sequences.
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                                                                                  Plant chromosome compositions and methods
Patent: WO 0055325-A 28 21-SEP-2000;
The University of Chicago (US)
Location/Qualifiers
                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Arabidopsis thaliana
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/db_xref="taxon:32630"
/note="Nucleic Acid"
6 c 3 g 5 t
                                     /db_xref="taxon:3702"
4 c 5 g
                                                           organism="Arabidopsis thaliana"
                                                                                                                                   Copenhaver, G. and Keith, K.
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Mammalia; Eutheria;
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Location/Qualifiers
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                                                                                                                                                                             Barany, F., Zirvi, W., Gerry, N.P., Favis, R. and Kliman, R. Method of designing addressable array for detection of sequence differences using ligase detection reaction Patent: WO 0179548-A 3236 25-OCT-2001, CORNELL RESEARCH FOUNDATION, INC. (US)
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/note="Hypothetical Probe Sequence"
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/db_xref="taxon:9606"
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Hammond,G.L. and Avvakumov,G.V.
Bammond,G.L. and Avvakumov,G.V.
DNA encoding variants of human corticosteroid
Patent: US 5432080-A 4 11-JUL-1995;
                                                                                  Sequence 4 from patent 134228 134228.1 GI:1825019
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1 (bases 1 to 21)
Hammond,G.L. and Avvakumov,G.V.
Variants of human corticosteroid binding globulin
Patent: US 5555969-A 4 21-JAN-1997;
Location/Qualifiers
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A BAC-based STS-content map spanning a 35-Mb region of hum chromosome 1p35-p36 Genomics 74 (1), 55-70 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (04-AUG-2001) Akira Horii, Tohoku University School Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Senda Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Pax:81-22-717-8047)
Location/Qualifiers
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Pred. No. 1.8e+05;
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                                                                                                                                                                                                                               Score 11.6; DI
Pred. No. 1.8e
0; Mismatches
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STS 8t8-AA040365
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Gunderson, K.

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AX116728
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                                                    1 (bases 1 to 21)
Goddijn,O.J. and Ohl,S.A.
PLASMIDS FOR PLANT TRANSFORMATION AND METHOD PACENT: WO 9901563-A 1 14-JAN-1999;
GODDIJN OSCAR JOHANNA MARIA (NL); OHL STEPHAN
LOCATION/QUALIFIERS
                                                                                                                                                                                          21 bp
Sequence 1 from Patent WO9901563.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     synthetic construct artificial sequences.
1 (bases 1 to 25)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 1851 26-APR-2001; Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1851 from Patent AX116728
                                                                                                                                             unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          synthetic construct.
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                                                                                                                                                           unidentified.
                                                                                                                                                                                 AX001122.1 GI:7241321
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                                                                                                                                  unclassified.
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/db xref="taxon:32630"
/note="Computer Generated Probe
5 c 7 g 4 t
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/db_xref="taxon:32630"
/note="Primer"
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                /db_xref="taxon:32644"
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                                  organism="unidentified"
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77.8%;
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Pred. No. 1.9e+05;
0; Mismatches 4
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Pred. No. 1
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AX153966
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DEFINITION
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AX175376
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A32924
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AX153966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human single nucleotide polymorphisms
Patent: WO 0138576-A 64 31-MAY-2001,
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 21)
Cargill,M., Ireland,J.S. and Lander,E.S.
                                                                                                                                                                                                              artificial sequences.
Location/Qualifiers
                                                                                                                                                                                                                                     synthetic construct
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           53
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/db_xref="taxon:32630"
8 c 4 g
                                                                                                                                                                                                                                                                      GI:1926573
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/db_xref="taxon:9606"
2 c 5 g 1
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            24 bp
from Patent WO0142308.
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80.0%;
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Pred. No. 2.4e+05;
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AR197424/c
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Sequence 509
AR090389
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Barbas, C.F. and Steinberger, P.
Barbas, C.F. and compositions useful for inhibiting infection of cells by hiv-1 patent: WO 0142308-A 53 14-JUN-2001; Novartis AG (CH); The Scripps Research Institutional Location/Qualifiers
             1 (bases 1 to 25)
Chenchik,A., Jokhadze,
Methods of assaying di
Patent: US 6352829-A
                                                                                                                     AR197424
Sequence 509 from patent
AR197424
                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 25)
Chenchik, A., Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 5994076-A 509 30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AX175376.1
                                                                         Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                Unclassified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AR090389.1
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k,A., Jokhadze,G. and Bibilashvilli,R. of assaying differential expression US 6352829-A 509 05-MAR-2002; Location/Qualifiers
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9 c 4 g
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|db_xref="taxon:32630"
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7 c 8 g
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Pred. No. 2.4e+05;
0; Mismatches 1;
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Pred. No. 2.4e+05;
0; Mismatches 1.
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TAKEDA CHEMICAL INDUSTRIES
OS Artificial Sequence
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             AX215063
Sequence 505 from Patent
AX215063
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Patent: JP 2001029090-A 44
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C12N1/15,C12N1/19,C12N1/21,C12N5/10,C12P21/08,(C12P21/02,
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/db_xref="taxon:32630"
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Patent: WO 0159103-A 505 16-AUG-2001;
RIBOZYME PHARMACEUTICALS, INC. (US); Blatt,
McSwiggen, James (US); Chowrira, Bharat M. (
Location/Qualifiers
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Sequence 1323 from Patent
AX422987
AX422987.1 GI:21526369
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1 (bases 1 to 17)
Blatt, L., Mcswiggen, J. and Chowrira, B.M.
Method and reagent for the modulation an
                                                                                  19 bp
Sequence 18 from Patent WO0129217.
AXII9996
AXII9996.1 GI:14036730
                                                                                                                                                                                                                                                                                                                                                                        Method and reagent for the inhibition of erg patent: WO 0188124-A 1323 22-NOV-2001; RIBOZYME PHARMACEUTICALS, INC. (US); GLAXO GROUP Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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artificial sequences.

1 (bases 1 to 19)
Prayaga, S. K., Taughier, R. J. and Bandaru, R.
Polypeptides and polynucleotides encoding
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Curagen Corporation (US)
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Scholar, E.M. and Iversen, P.L.
Phosphorothicate inhibitors of metastatic breast cancer
Patent: US 5552390-A 9 03-SEP-1996;
Location/Qualifiers
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Sequence 9 from patent US 5552390.
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6: /cgn2_6/ptodata/2/ina/backfiles1.seq;*
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US-09-593-012-191
US-09-593-012-293
US-09-593-012-213
US-09-593-012-24
US-08-233-608-24
US-08-233-012-24
US-08-233-012-24
US-08-233-012-12-24
US-09-593-012-12-24
US-09-593-012-19-70
US-09-593-012-19-70
US-08-887-480-64
US-08-887-480-64
US-08-08-21-293-812-64
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CURRENT FILING DATE: 2000-06-13
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PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
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US-09-593-012-191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Trichoderma harzianum
US-09-593-012-195
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                                                                                                                                               Sequence 191, Application US/09593012 Patent No. 6387652 GENERAL INFORMATION:
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Best Local
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GENERAL INFORMATION:
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               APPLICANT: HAUGLAND, Richard
APPLICANT: VESPER, Stephen
TITLE OF INCENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND
FILE REFERENCE: HAUGLAND-1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
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APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
FILE REFERENCE: HAUGLAND=1A

APPLICATION NUMBER: US
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US-09-428-219-11
US-09-428-219-18
US-09-280-805-34
US-09-280-805-51
US-09-280-805-159-30
US-09-101-886B-15
US-09-290-640-27
US-09-117-090-13
US-09-113-932-128
US-09-313-932-128
US-09-313-930-12
US-09-313-930-13
US-09-313-930-13
US-09-417-082-10
US-09-417-082-10
US-09-417-082-10
US-09-417-927-11
US-09-117-927-11
US-09-117-927-11
  09/290,990
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Pred. No. 4.1e+02;
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FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR PILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
SOFTWARE: Patentin version 3.0
SEQ ID NO 203
LENGTH: 21
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; TYPE: DNA
; ORGANISM: Trichoderma asperellum/hamatum
US-09-593-012-191
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US-09-593-012-194
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US-09-593-012-203
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PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 194
LENGTH: 21
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APPLICANT: HAUGLAND, Richard
APPLICANT: VESPER, Stephen
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Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
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NUMBER OF SEQ ID NOS: 225
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PRIOR APPLICATION NUMBER: US 60/081,773
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                                                                                                                                                                                                                                                             APPLICANT: HAUGLAND, APPLICANT: VESPER, 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Trichoderma asperellum/hamatum/viride
ORGANISM: Trichoderma viride/atroviride/koningii
                                                                                                                                                                                                                                            PPLICANT: VESPER, Stephen IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
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Pred. No. 4.2e+02;
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Pred. No. 4.
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US-09-593-012-31
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SEQ ID NO 31
LENGTH: 22
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Best Local S
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NUMBER OF SEQ ID NOS: 225
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CURRENT FILING DATE: 2000-06-13
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APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
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                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                               ent No.
              ATTORNEY/AGENT INFORMATION:
NAME: Spruill, W. Murray
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                          PPLICANT: Ligon, James P
                                                                                                                                                                                                                                   CITY: Hawthorne STATE: NY
                                                                                                                                                                                                                                                                                                                                  TLE OF INVENTION: Detection of Fungal Pathogens Using the
                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                    COUNTRY: U
REFERENCE/DOCKET NUMBER: CG
                                                                                          APPLICATION NUMBER:
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                                                             ASSIFICATION:
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7 Skyline Drive
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CGC 1739
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Pred. No. 4.2e+02;
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Pred. No. 4.
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HYPOTHETICAL:
ANTI-SENSE:
US-08-887-480-24
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US-08-887-480-24
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Query Match
Best Local Similarity
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 20 base pairs
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                    MOLECULE TYPE: Other nucleic acid
MOLECULE TYPE: Other nucleic acid
DESCRIPTION: Oligonucleotide primer JB548
                                                                                                                                                                                                                               NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE DOCKET NUMBER: CG
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/887,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 15-OCT-1996
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DESCRIPTION:
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TELEPHONE: 919-541-8615
                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 10591
                                                                                                                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                         nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tarrytown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: No. 5814453artis Corporation 520 White Plains Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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Oligonucleotide primer JB548
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82.4%;
 71.8%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymerase Chain Reaction
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                                                                                                                                                                                                                                                                                                                           US 08/722,187
                                                                                                                                                                                         24:
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 Score
Pred.
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Pred. No. 5
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 12.2; DB 1;
No. 5.4e+02;
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RESÚLT 9
PCT-US95-04712-24
                    Sequence 24, Application PC/TUS9504712
GENERAL INFORMATION:
APPLICANT: Ligon, James M
APPLICANT: Beck, James J
TITLE OF INVENTION: Detection of FA
TITLE OF INVENTION: Polymerase Chai
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Best Local S
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INFORMATION FOR SEQ ID NO: 24:
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                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: C
DESCRIPTION: (
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA: US 08
APPLICATION UNMBER: US 08
FILING DATE: 04-APR-1994
PILING DATE: 1NFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIDIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TILE OF INVENTION: Detection of Fungal Pathogens Using the TILE OF INVENTION: Polymerase Chain Reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Walsh, Andrea C.
REGISTRATION NUMBER: 34,988
REFERENCE/DOCKET NUMBER: CGC 1739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             t: 20 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                               Similarity
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7 Skyline Drive
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Beck, James J
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Oligonucleotide primer JB548
                      Detection of Fungal Pathogens Using the Polymerase Chain Reaction
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Pred. No. 5.4e+02;
0; Mismatches
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; ANTI-SENSE: NO
PCT-US95-04712-24
                                                                          ; ORGANISM: Penicillium brevicompactum/alberechii US-09-593-012-120
                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-593-012-120
Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                  SOFTWARE: PatentIn version 3.0 SEQ ID NO 120
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 120, Application US/09593012 Patent No. 6387652
                                                                                                                                                                                                   FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 225
                                                                                                                                                                                                                                                                                                                                   APPLICANT: VESPER, Stephen
FITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTER:
                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Walsh, Andrea C. 988 REGISTRATION NUMBER: 34,988 REFERENCE/DOCKET NUMBER: CG TELECOMMUNICATION INFORMATION: TELEPHONE: 919-541-8666
                                                                                                                                 LENGTH:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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     70.6%; Score 12; DB 4; Liarity 100.0%; Pred. No. 6.8e+02; Conservative 0; Mismatches 0;
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Pred. No. 5.4e+02;
0; Mismatches 3;
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                                       Length 17;
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         Indels
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US-09-428-219-70/c
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SEQ ID NO 70
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 70, Application US//
Patent No. 6177273
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -08-291-932A-812/c
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CURRENT FILING DATE: 1999-10-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ANTISENSE MODULATION OF INTEGRIN-LINKED KINASE EXPRESSION FILE REPERENCE: RTS-0101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                               OPERATING SYSTEM: IBM P.C.
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/2
                                               PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                              ITLE OF INVENTION:
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COUNTRY:
                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
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                                                                                                    PILING DATE: AN CLASSIFICATION:
                                                                                                                                                                                                  COMPUTER:
   FILING DATE: May 1
APPLICATION NUMBER
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                                   APPLICATION NUMBER:
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  McSwiggen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stinchcomb, Dan T.
Draper, Kenneth G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                              IBM Compatible
May 18, 11,
                                                                                                              August 15, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                   RELATED TO LEVELS OF NF-KB
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DISEASES OR CONDITIONS
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                                  08/245,466
                                                                                                                             US/08/291,932A
                                                 including application described below:
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Sequence 57, Apprince 57, Apprince 57, Apprince 5814453
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Best Local Similarity
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                                                                                             TELEPAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 16 base pairs
                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
RAPPLICATION UMMBER: US 08/722
FILING DATE: 15-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meidg. J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: CGC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587
                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
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                                  MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ITLE OF INVENTION: .Detection of Fungal Pathogens Using the ITLE OF INVENTION: Polymerase Chain Reaction
                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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                                                               TRANDEDNESS:
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                                                                               H: 16 base pairs
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/desc = JB570"
                               other nucleic acid
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92.3%;
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                 "Oligonucleotide primer
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Pred. No. 1
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Query Match

67.1%;

Score 11.4;

DB 1;

Length 16;

NUMBER OF SEQUENCES:

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US-08-722-187-57
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Patent No. 5814453
                                                                         Sequence 57, App...
No. 595527
                                                                                                                                                                                                                                                     Matches
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                                                              Patent No. 5955274
GENERAL INFORMATION:
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TITLE OF INVENTION: Detection of Fungal Pathogens Using the
TITLE OF INVENTION: Polymerase Chain Reaction
NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/887,480
PILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/722,187
PILING DATE: 15-OCT-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timochy
REGISTRATION NUMBER: 38,241
REPERENCE/DOCKET NUMBER: CGC 1739/P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                  DESCRIPTION:
HYPOTHETICAL:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Oligonucleotide primer
DESCRIPTION: JB577"
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APPLICANT: Ligon, James
APPLICANT: Beck, James
TITLS OF INVENTION: Det
TITLS OF INVENTION: Pol
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TELEPHONE: 919-541-8587
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12; Conserv
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                                                                                                 Application US/08722187
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                                                James M
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Detection of Fungal Pathogens Using the Polymerase Chain Reaction
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Pred. No. 1.4e
0; Mismatches
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0; Mismatches 1
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Best Local Similarity
Thes 12; Conserve
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HYPOTHETICAL:
US-08-722-187-57
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US-08-722-187-64
                                                                                                                                                                                                                                                                                                                   Sequence 64, App. 8955
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: 111
MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US 0:
FILING DATE: 04-APR-1994
ATTORNEY/AGENT INFORMATION:
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,187 FILING DATE:
                                                                                   ZIP: 10532
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: CG
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                              NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                            APPLICANT: Beck, James J
TITLE OF INVENTION: Detection of Fungal Pathogens Using the
TITLE OF INVENTION: Polymerase Chain Reaction
                                                                                                                                                                                                                                                                                       APPLICANT:
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STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Walsh, Andrea REGISTRATION NUMBER:
                                                                                                                                                                                ADDRESSEE:
STREET: 7
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                                                                                                                                                                  CITY: Hawthorne
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/desc = "Oligonucleotide primer
JB570"
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92.3%;
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Pred. No. 1.4e+03
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RESULT 18
PCT-US95-04712-57
; Sequence 57, Application PC/TUS9504712
; GENERAL INFORMATION:
; GENERAL INFORMATION:
APPLICANT: Ligon, James M
; APPLICANT: Beck, James J
; TITLE OF INVENTION: Detection of Fungal Pathogens Using the TITLE OF INVENTION: Polymerase Chain Reaction
; TITLE OF INVENTION: Polymerase Chain Reaction
                                                                                                                                                                                                                               문
                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Trichoderma longibrachiatum/citreoviride
US-09-593-012-198
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 198
                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HAUGLAND, APPLICANT: VESPER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Oligonucleotide primer
DESCRIPTION: JB577"
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APPLICATION NUMBER: US 0
FILING DATE: 04-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                198,
                                                                                                                                                                                                                                                                                                 12,
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92.3%;
                                                                                                                                                                                                                                                                                          67.1%; Score 11.4; DB 4; Length 16; 92.3%; Pred. No. 1.4e+03; cive 0; Mismatches 1; Indels
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Pred. No. 1
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PCT-US95-04712-57
ZIP: 10532

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION UMBER: PCT/US95/04712

FILING DATE:

CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                 Sequence 64. Application PC/TUS9504712
GENERAL INFORMATION:
APPLICANT: Ligon, James M
APPLICANT: Beck, James J
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Best Local S
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/233,608
APPLICATION UNMBER: 194
ATTORNEY/AGENT INFORMATION:
NAME: Walsh, Andrea C.
REGISTRATION NUMBER: 34,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 16 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear nucleic acid MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "Oligonucleotide primer DESCRIPTION: /B570"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: CG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Ciba-Geigy Corporation
STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                           IIILE OF INVENTION: Detection of Fungal Pathogens Using the
IIILE OF INVENTION: Polymerase Chain Reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                STATE:
                                                                                                                                                                                                                                                     ADDRESSEE: Ciba-Geigy Corporation STREET: 7 Skyline Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US95/04712 FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: NY
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92.3%;
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Pred. No. 1.4e+03;
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US 08/233,508

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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 16 base pairs
                                 TELEFAX: 212-218-2200
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                   OPERATING SYSTEM: MS-DOS Ver3.30
SOPTWARE: PATENT AID Ver1.0
CURRENT APPLICATION DATA:
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                                                                                                                                                               APPLICATION NUMBER: US/08/606,505B FILING DATE: 23-MAR-1996 PRIOR APPLICATION DATA: APPLICATION NUMBER: JP44963/92 FILING DATE: 02-MAR-1992
                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.50 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: other nucleic acid
                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Perry, Lawrence S.
REGISTRATION NUMBER: 31865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 212-218-2100
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Local Similarity 92.3%;
les 12; Conservative
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                 ENGTH:
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E: New York
TRY: U.S.A.
10112-3801
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nucleic acid
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SHIMADA, Yukihisa
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OKINAKA, Yasushi
JENTION: NOVEL PLANT GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OHBAYASHI, Masaya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KIKUCHI, Yasuhiro
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JB577"
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Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                                                                                                       720 Kb storage
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RESULT 22 US-08-652-127C-1

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; Sequence 1, Application US/08652127C

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TOPOLOGY: linear

MOLECULE TYPE: Other nucleic acid

DESCRIPTION: Synthetic DNA

SEQUENCE DESCRIPTION: SEQ ID NO: 19
US-09-616-990-19
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US-09-616-990-19/c
; Sequence 19, Application US/09616990
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US-05-606-505B-19
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Best Local Similarity
Matches 12; Conserv
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GENERAL INFORMATION:
                                                                  Matches
                                                                                                Query Match
                                                                                Best Local
                                                                                                                                                                                                                                                                                          INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.50 inch,
COMPUTER: IBM PS/V
OPERATING SYSTEM: MS-DOS Ver3.30
SOFTWARE: PATENT AID Ver1.0
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/616,990
FILING DATE: 14-Jul-2000
PRIOR APPLICATION DATA:
APPLICATION UMBER: JP44963/92
FILING DATE: 02-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: KIKUCHI, Yasuhiro
KIYOKAWA, Shigeto
SHIMADA, Yukihisa
                                                                                                                                                                                                                                                                                                                                        NAME: Perry, Lawrence S.
REGISTRATION NUMBER: 31865
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OKINAKA, Yasushi
TITLE OF INVENTION: NOVEL PLANT GENES
                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18 TGCTTNGGNGGGATC
                                 3 TGCTTCGGCGGGAAC 17
                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEB: FITZPATRICK, CELLA, HARPER & SCINTO STREET: 30 Rockefeller Plaza
                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10112-380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
                                                                                                                                                                                                                                        LENGTH: 18 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                        TELEFAX: 212-218-2200
N FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                          TELEPHONE: 212-218-2100
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Synthetic TMA
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                                                                67.1%; Score 11.4; DB 4; Length 18; 80.0%; Pred. No. 1.4e+03; Indels
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           720 Kb storage
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Patent No. 5792611
GENERAL INFORMATION:

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RESULT 23
US-08-358-556A-1
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                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Chatelain, FrancAPPLICANT: Kumarev, Viktor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (613) 232-5831 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                              TITLE OF INVENTION: ITTLE OF INVENTION: ITTLE OF INVENTION: NUMBER OF SEQUENCES:
                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compactible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: ASCII
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE PORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Jacobson, Price, Holman & Stern
STREET: 400 Seventh St. N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: George A. Seaby
REGISTRATION NUMBER: 24,034
REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
TELEPHONE: (613) 232-5815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08, FILING DATE: May 23, 1996 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
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TITLE OF INVENTION: DETECTION OF PLANT
TITLE OF INVENTION: PATHOGEN FUNGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TUMBER OF SEQUENCES:
                                                                                                                                                           COUNTRY:
                                                                                                                                                                              CITY: Washington D.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       880 Wellington Street, Suite 708
                                                                                                                                                           U.S.A.
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Seaby & MacLean
14-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                              Process for Preparing Polynucleotides on a Solid Support and Apparatus Permitting its Implementation
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              US/08/358,556A
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US-08-358-556A-1
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Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6, Apr-
Sequence 6, Apr-
No. 6033909
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                          CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/003,972
FILING DATE: 19-JAN-1993
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL: NO ANTI-SENSE: NO FRAGMENT TYPE: 1
                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IHM PC compacible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FR 9: FILING DATE: 16-DEC-1993 ATTORNEY/AGENT INFORMATION:
                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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LENGTH: 18 base pairs
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CELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        IITLE OF INVENTION: Oligonucleotide IITLE OF INVENTION: Preparation and NUBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
             NAME: Einaudi, Carol P.
REGISTRATION NUMBER: 32,220
REFERENCE/DOCKET NUMBER: 02
                                                                                                                                                         APPLICATION NUMBER: US/08/281,203 FILING DATE: 27-JULY-1994
                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                       CITY: Washington
                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Finnegan, Henderson, Farabow, Garrett & ADDRESSEE: Dunner
STREET: 1300 I Street, N.W.
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81.2%;
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Pred. No. 1
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Use
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                                                                                                                                                                                                                                                                                                      US-08-005-283-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 08/777,266
PRIOR FILING DATE: 1996-12-31
NUMBER OF SEQ ID NOS: 226
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 146
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6319906
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local
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                                                                                                                                                                                                                                                                     Patent No. 564626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: ISPH-0376
CURRENT APPLICATION NUMBER: US/09/326,1868
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Vickers, Timothy A.
TITLE OF INVENTION: Oligonucleotide Compositions and Methods for the TITLE OF INVENTION: Modulation of the Expression of B7 Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bennett, Clarence Frank
APPLICANT: Vickers. Timorby a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                       APPLICANT: Helsberg, Matthias
APPLICANT: Winkler, Irvin
TITLE OF INVENTION: 3'-Derivatized Oligonuclectide Analogs
TITLE OF INVENTION: 3'-Derivatized Oligonuclectidic Groupings, Their Preparation and
TITLE OF INVENTION: with No. 5646261-Nuclectidic Groupings, Their Preparation and
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
                          STREET: 1300 1 50
CITY: Washington
                                                             ADDRESSEE:
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            146, Application US/09326186B
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                                                                                                                                                                                                                                                     INFORMATION:
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                                                  E: Dunner
1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                  O'Malley, Gerard
Helsberg, Matthias
                                                                                                                                                                                                                   Uhlmann, Eugen
Peyman, Anuschirwan
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                                                                              Finnegan, Henderson, Farabow, Garrett
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.9%; Score 11.2;
81.2%; Pred. No. 1.
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pred. No. 1.
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                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATION
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Pacentin Release #1.0, Version #1.25
SOFTWARE: Pacentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNUMBER: US/08/005,283
FILING DATE: 19-JAN-1993
CLASSIFICATION: 536
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: DE P42
APPLICATION UNMBER: DE P42
FILING DATE: 22-JAN-1992
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: TBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin. Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 02:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: 3'-Derivatized Oligonucleotide Analogs
TITLE OF INVENTION: with No. 5646261-Nucleotidic Groupings, Their Preparation
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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              FILING DATE: 22-JAN-1992
ATTORNEY/AGENT INFORMATION:
                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 202-408-4400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear DNA
                                                                                                                                                                                                                                                                                              STREET: 1300 1 STREET: Washington
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                                                                                                                                                                                                                                                           COUNTRY:
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13; Conserv
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20005-3315
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Hammond, Alan W.
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1300 I Street, N.W.
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Helsberg, Matthias
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81.2%; Pred. No. 1.
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                                                       DE P4201663.0
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RESULT 29
US-08-472-255A-56
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US-09-593-012-48/c
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SEQ ID NO 48
LENGTH: 18
TYPE: DNA
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Patent No. 6387652
GENERAL INFORMATION:
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Matches 13, Conservative
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APPLICANT: VESPER, Stephen
TITLE OP INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR PILING DATE: 1999-04-14
PRIOR PILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
NUMBER OF SEQ ID NOS: 225
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                          ENERAL INFORMATION:
                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                               TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS TITLE OF INVENTION: TO SELECTINS (AS AMENDED)
                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                       APPLICANT: PARMA, DAVID
APPLICANT: GOLD, LARRY
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TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
OMPUTER READABLE FORM:
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REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION:
                             COUNTRY:
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                                                                                                    ADDRESSEE:
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TYPE: nucleic acid
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                                                                             3: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite 200
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100.0%;
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81.2%; Fred. No. 1.9e+03
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; OTHER INFORMATION:
US-08-472-255A-56
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Best Local Similarity
                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: PARMA, DAVID
APPLICANT: GOLD, LARRY
TITLE OF INVENTION: HIGH APPINITY NUCLEIC ACID LIGANDS
TITLE OF INVENTION: TO LECTINS
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                              COMPUTER: IBM pc compatible OPERATING SYSTEM: MS-DOS SOPTWARE: WordPerfect 6.0 CURRENT APPLICATION DATA:
                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/714,131
PILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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FILING DATE: 21-OCTOBER-1992
ATTORNEY/AGENT INFORMATION:
NAME: BAITY J. SWAIDON
REGISTRATION NUMBER: 33,215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/53
FILING DATE: 11-JUNE-1991
PRIOR APPLICATION DATA:
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                FILING DATE: 0 CLASSIFICATION:
                             APPLICATION NUMBER: US/08/479,724A FILING DATE: 07-JUNE-1995
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                                                                                                                                                                                              COUNTRY:
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LECOMMUNICATION INFORMATION:
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00 E. Prentice Avenue, Suite 200
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                                                                                                                                   Diskette, 3 1/2 diskette, 1.44 MB
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Pred. No. 2.4e+03;
1; Mismatches · 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tent No.
                                                                                                                                 SOFTWARE: WordPerfect 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,256B
FILING DATE: 07-JUNE-1995
                                                                                                                                                                                                                                                                              STREET: BYVC
CITY: Englewood
CTATE: Colorado
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            PILING DATE: 11-JUNE-1991 PRIOR APPLICATION DATA:
                                                        APPLICATION NUMBER: 07/714,131
FILING DATE: 10-JUNE-1991
PRIOR APPLICATION DATA:
                                                                                                    CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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REFERENCE/DOCKET NUMBER: NEX40-1
REFERENCE/DOCKET NUMBER: NEX40-1
REFERENCE/DOCKET NUMBER: 33,333
TET.RPHONE: (303) 793-3333
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APPLICATION NUMBER:
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                        TLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS TLE OF INVENTION: TO LECTINS MBER OF SEQUENCES: 177
                                                                                                                                                                                                                                                                            COUNTRY:
                                          APPLICATION NUMBER:
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PPLICATION NUMBER:
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8400 E. Prentice Avenue, Suite 200
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E: RNA
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                                        07/536,428
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07/964,624
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                                                                                                                                                                                                                                3 1/2 diskette, 1.44 MB
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-952-793-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION: APPLICANT: PARMA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION:
                                                       APPLICATION NUMBER: 08/47 FILING DATE: 07-JUNE-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                   FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
                                                                                                                                                 FILING DATE: 07-JUNE-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM DC compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordberfect 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/472,256
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APPLICATION NUMBER:
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ELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                            APPLICATION NUMBER: 08/47)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 8400 E. CITY: Englewood
         REFERENCE/DOCKET NUMBER: NEX40C/PCT
                                                                                                                                                                                                                                                                 FILING DATE: 05-JUNE-1995
                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                 APPLICATION NUMBER:
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                                        Barry J. Swanson
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00 E. Prentice Avenue, Suite
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                                                                                                                                    08/472,255
                                                                                      08/477,829
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Pred. No.
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US-08-952-793-56
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Best Local Similarity 90.9
Matches 10; Conservative
TELEPAX: (303, INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS: SEQUENCE CHARACTERISTICS: Dase pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -US96-09455A-56
                                                          NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEPAX: (303) 793-3433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             quence 56, App.
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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: RNA
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MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: 08/472,256
FILING DATE: 07-JUNE-1995
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FILING DATE: 07-JUNE-1995
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                                                                                                                                                       LICATION NUMBER: 08/477,829
ING DATE: 07-JUNE-1995
NEY/AGENT INFORMATION:
                                                                                                                                                                                                      APPLICATION DATA:
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                                                                                                                                                                                                                  CATION NUMBER: 08/47
                                                                                                                                                                                                                                                                                                                                                                                                                   WARE: WordPerfect 6.0 PAPPLICATION DATA:
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Colorado
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PARMA, et al.
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O E. Prentice Avenue, Suite 200
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                                                                                                            NEX40C/PCT
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Pred. No.
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Best Local Similarity 90.5
Matches 10; Conservative
                                                                                                                                                                                                                     FILING CLASSIFICATION: CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 60/001,135
APPLICATION NUMBER: 60/001,735
TYPE DATE: July 13, 1995
TYPE DATE: July 13, 1995
TYPE DATE: 71978R: 08/300,726
                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: RNA
FEATURE:
OTHER INFORMATION:
FEATURE:
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                                                                                                                  REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/0
FILING DATE: July 12, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                    FILING DATE: September 2, 1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
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TITLE OF INVENTION:
                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 90071-2066
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Los Angeles
California
                nucleic acid
                                                                                                                                                                                      Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         633 West Fifth Street
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                                                                                                      (213)
                                                                                                                                                                                                                                                                                                                                                                                Word Perfect 5.1
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Merlo, Donald J.
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single
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Pred. No. 2.4e+03;
1; Mismatches 0; Indels
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Matches Query Notels Best Local

Similarity

64.5%; 85.7%;

Score 10.8; DB Pred. No. 3e+03; Mismatches

4; Length 18;

Conservative

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Indels . 0;

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US-08-265-429A-2
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                                                                                                                                                                                                                 US-09-156-856-11/c
                                                                                                                                                                                                                                                        RESULT 36
                                                                                                              Sequence 11, Application US/09156856A Patent No. 6221591 GENERAL INFORMATION:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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APPLICANT: Aerts, Johannes M.
TITUE OF INVENTION: Determination of a genetic risk factor for infection
TITUE OF INVENTION: and other diseases, and detection of activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQUENCE CHARACTERISTICS:
LENGTH: 20 nucleon:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US
FILING DATE: 24/06/94
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; SOFTWARE: PatentIn Ver. 2.0; SEQ ID NO 11; LENGTH: 20; TYPE: DNA; ORGANISM: Homo sapiens US-09-156-856-11
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US-09-367-007C-14/c
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US-09-659-791A-59/c
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                             NUMBER OF SEQ ID NOS: 39
SEQ ID NO 14
SENGTH: 20
TYPE: DNA
ORGANISM: artificial sequence
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Matches
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CURRENT FILING DATE: 2000-09-11
NUMBER OF SEQ ID NOS: 90
SEQ ID NO 59
LENGTH: 20
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                                                                                                                                                                                                                                                                                             Sequence 14, Application US/09367007C Patent No. 6416987
                                                                                                                                                                                                                                                                                   GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/156,856A
CURRENT FILING DATE: 1998-09-18
NUMBER OF SEQ ID NOS: 20
                                                                                                             FILE REFERENCE: D5978
CURRENT APPLICATION NUMBER: US/09/367,007C
CURRENT FILING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: PCT/US98/02145
PRIOR FILING DATE: 1998-01-03
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                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                              APPLICANT: Bertino, Joseph R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
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                    FEATURE:
                                                                                                                                                                                               ITLE OF INVENTION: Mutants of Thymidylate Synthase and Uses Thereof
NAME/KEY: primer_bind
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12; Conserv
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                                                                                                                                                                                                                  Liu-Chen,
                                                                                                                                                                                                                                Tong,
                                                                                                                                                                                                                                            Banerjee, Debabrata
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Pred. No. 3.1e+03;
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PCT-US95-09069-2
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                                                                     Query Match
Best Local
                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 822..803
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                               NAME: Nelson, M. Bud
REGISTRATION NUMBER: 35,300
REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                      DRIGINAL SOURCE:
                                                                                                                                                                                                   OLECULE TYPE:
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                                                                                                                                            STRAIN: YX36
                                                                                                                                                        ORGANISM:
                                                                                                                                                                                                                                                                                                                      TELEPHONE:
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                                                                                                                                                                                                                               STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION: Antisense primer hTS-6B for PCR amplification of part INFORMATION: of 890 base pair fragment of human TS gene INFORMATION: from nucleotide 50 to the C-terminus
                                                                                                                                                                                                                     POLOGY:
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NT APPLICATION DATA:
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                                                    12; Conservative
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14203-2391
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                                                                                                                                                                                                                                              nucleic acid
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    Hodgson, Russ, Andrews, Woods & Goodyear
1800 One M&T Plaza

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                                                                                                                             bacterium
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                                                                                                                                                                                                                              single-stranded
                                                                                                                                                                                     synthesized
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                                                                    63.5%;
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                                                    Score 10.8; D
Pred. No. 3.1e
0; Mismatches
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Pred. No. 3.1e
0; Mismatches
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                                                                  3.1e+03;
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les 2;
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US-08-974-549A-510
                                                                                                                                                          TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                         NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER
FILING TO DATA: 14-AUG-1997
APPLICATION NUMBER
FILING TO THE PRIOR NUMBER
FILING TO THE PRIOR NUMBER NU
                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
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(USSI) US ENVIRONMENTAL PROTECTION AGENCY

Claim 62; Page

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13-JUN-2000; 2000US-0593012

14-MAY-2002.

Stachybotrys chartarum

ulcer; asthma; allergy; sinusitis; PCR; primer; ss.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to specific fungi or bacteria detected and CC quantified in a sample by extracting and recovering DNA from the organism CC in the sample and hybridising and amplifying the DNA sequence using CC sequences unique to the species/group of species but common to all CC isolates of the species/group of species. The method especially employs Sequences (ABN99429-ABN99653) for detection of specified fungi/fungal CC groups or specified bacteria. The method enables detection and CC quantification of specific fungi or bacterial primer and treatment of CC fungal/bacterial species, useful medically for disgnosis and treatment of CC fungi or bacteria associated with health problems such as infections, CC gastroenteritis, ulcers, asthma, allergies and sinusitis. It is also CC useful to detect and/or quantify microorganisms in the environment e.g. CC to establish the risk of adverse health effects (e.g. pulmonary CC haemorrhage from levels of Stachybotrys chartarum in air samples in
                                                                      Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe
                                                       analysis
                                                                                                                                                                                                                                                            Haugland R,
                                                                                                                                                                                                                                                                                                                                                                                 13-JUN-2000; 2000US-0593012
                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUN-2001; 2001WO-US18892.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stachybotrys chartarum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Stachybotrys chartarum reverse PCR primer
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100.0%; Pr
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Pred. No. 2.9e+02;
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RESULT 3
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ORGANISM
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  Query Match
Best Local Similarity
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ307451 21 bp DNA linear GSS 29-SEP-20 1M0009E09F Mouse 10kb plasmid UUGCIM library Mus musculus genomic Clone UUGCIM0009E09 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0009 row: E column: 09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         house mouse.
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                                                                           was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hint end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114 [gb]AF129072.1], a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent coli X110-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone_lib="Mouse 10kb plasmid UUGC1M library"
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|strain="C57BL/6J"
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lone="UUGC1M0009E09"
58.8%;
100.0%;
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  Score 10;
Pred. No.
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  DB 17; Length 21; 3.8e+05;
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RESULT 5
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AI421881/c
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bonaldo, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AI421881 25 bp mRNA linear EST 30-MAR-1999 tf55h03.x1 NCI CGAP Brn23 Homo sapiens cDNA clone IMAGE:2103221 3' similar to TR:P93237 P93237 PROLINE-RICH PROTEIN PRP2 PRECURSOR. ;,
                                                                                                                           N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primatee; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 25)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index (1998)
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Location/Qualifiers
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Seg primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AI421881.1 GI:4267812
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                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                  TICCTIGGGCGGG 4
                                                                                                                              TIGCTICGGCGGG 14
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                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                              adaptors (Pharmacia), digested with Not I at the Not I and Eco RI sites of the modified Library is normalized, and was constructed Soares and M.Fatima Bonaldo."

13 c 3 g 1 t
                                                                                                                                                                                                                                                                                                                                                                        T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and clor
                                                                                                                                                                                                                                                                                                                                                                                                                    /db xref="taxon:9606"
/clone="IMARS:2103221"
/clone lib="NCI CGAP Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Homo sapiens"
                                                                                                                                                                                         57.6%;
84.6%;
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                                                                                                                                                                       Score 9.8; DB 9;
Pred. No. 4.9e+05;
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Sequencing Center information can be

M. Fatima

I and cloned into

pT7T3 vector. by Bento

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RESULT 5
AAZ59948
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The invention relates the identification of genes whose expression CC levels are altered by chronic exposure to one or more drugs of abuse CC (e.g., ethanol, stimulants, opiates). The methods of the invention CC monitor the response of a cell to a drug of abuse, and comprise CC contacting the cell with the drug of abuse, and detecting the expression of the contacting the cell with the drug of abuse, and detecting the expression for CC probes that specifically hybridise to the ESTs. The methods are used to CC identify genes whose expression levels are altered ky chronic or acute whose regulation is altered in alcohol tolerance and/or addiction CC whose regulation is altered in alcohol tolerance and/or addiction CC useful in designing therapeutic and recovery regimens, and in evaluating the susceptibility of an organism or patient to drugs in a medical CC context. Monitoring the expression of identified genes and/or ESTs CC provides a mechanism by which test agens can be screened for the ability to alter or modulate the response of the organism to drugs of abuse. CC gramers used to amplify 4 cDNA hybridisation probes from SH-SY5Y-AH1861 cnmarroblastoma cell total RNA. The probes were used in Northern CC blot analysis of gene expression in control and ethanol-treated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                문
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21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Drug exposure; drug abuse; gene expression; EST; expressed sequence tag; identification; tolerance; addiction; ther screening, cellular response; ethanol; expression analysis; Northern blot; delta-like protein; DLK; reverse transcriptase-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human delta-like
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Page 69; 98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lai C,
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99US-0337022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SH-SY5Y-AH1861 cells in an exemplification of the present invention. The genes whose expression was analysed were dopamine beta-hydroxylase (DBH and sodium- dependent norepinephrine transporter (NTT), both of which a involved in norepinephrine metabolism; delta-like protein (DLK); and monocyte chemoattractant peptide 1 (MCP-1). These genes are thought to be important targets of ethanol, Primers AAZ59948-Z59949 were used to generate the delta-like protein (DLK) probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA; enzyme; enzymatic RNA molecule; ERM; cleave; RNA; mRNA; HnRNA; picornavirus; HIV; immunodeficiency virus; hepatitis B virus; HBV; papilloma virus; HPV; Epstein-Barr virus; EBV; TCLV; papilloma virus; hBV; Epsteititis C virus; HCV; cytomegalovirus; influenza virus; HSV; herpes simplex virus; vector; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ50799 standard;
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JOURNAL COMMENT
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AUTHORS
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Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0019 row: E column: 20
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Sedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R. and Wright, D., Weise, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 bp DNA linear GSS 16-FEB-200
2M0019E20R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0019E20 R, DNA sequence.
AZ781461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 GCTTCGGCGGGAAC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A2781461.1 GI:12914177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       house mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ass: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                            quality sequence stop: 19.
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/db_xref="taxon:5911"
/clone_lib="Chilcoat_Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713:"
7 c 6 g 4 t
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high mclar excess. The
                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Tetrahymena thermophila"
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                                                                                                                                                                                                                                                                      organism="Mus musculus"
strain="C57BL/6J"
db_xref="taxon:10090"
clone="UUGC2M0019E20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     strain="CU428.1"
                                                                                                                                      aboratory Mouse DNA Resource
                                                                                                                                                                                                                              sex="Male"
                                                                                                                                                                                                                                                clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                                                                                                                                                                                                                       ocation/Qualifiers
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78.6%;
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Pred. No. 9.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2
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KEYWORDS
SOURCE
ORGANISM
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AZ432681
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia, Eutheria, Rodentia, Sciurognathi, Murid 1 (bases 1 to 20)
Dunn,D., Aoyagi.A., Barber,M., Beacorn,T., Duval, Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Ped,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0218 row: N column: 11
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
High quality sequence stop: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nouse mouse.
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Laboratory Mouse NAM RESOURCE (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymerase and T4 polymeride were polymuclectide kinase. Adaptor oligonucleotides were polymuclectide with T4 DNA polymerase and T4 polymuclectides were polymuclectide with T4 DNA polymerase and T4 polymuclectide were polymuclectide with T4 DNA polymerase and T4 polymer
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                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F. note="Vector: PMP42NV; Purified genomic DNA from the Jack musculus C578L/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="UUGC1M0218N11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
strain="C57BL/6J"
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78.6%;
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                                                                                                                                                                                                                                                                                            Jackson
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Reilly
                                                                                                                                                                                                        The DNA
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14; 44pp; English

The present invention relates to a member of the family of proteins known CC as ADAMTS (A Disintegrin And Metalloproteas Thrombospondin domain) CC proteins, the new member being designated ADAMTS-SI, antibodies to polynucleotides encoding ADAMTS-SI, assays for determining CC assays for studying the function of ADAMTS-SI, assays for determining CC assays for studying the function of ADAMTS-SI, and to the use of ADAMTS-SI assays for determining CC apolypeptides or polynucleotides in diagnostic, biotherapeutic, or gene CC therapy methods. ADAMTS proteins and ADAMTS protein agonists and CC antagonists are useful in the manufacture of a medicament for treating CC asthmat, chronic obstructive pulmonary disease, Alzheimer's disease, organ transplant toxicity and rejection, cachexia, allergy, cancer (such as solid tumour cancer including colon, breast, lung, prostate, brain, and CC haematopoietic malignancies including leukaemia and lymphoma), tissue ulcerations, restenosis, periodical disease, epidermolysis bullosa, cateoporosis, loosening of artificial joint implants, atherosclerosis consensative heart failure, myocardial confirmation, stroke, cerebral ischemia, head trauma, spinal cord injury, contingent of disease, parkinson's disease, migraine, dapression, coular anglogenesis, corneal injury, macular degeneration, occupantion enhancement, amyotrophic lateral sclerosis, multiple sclerosis, coular anglogenesis, corneal injury, macular degeneration, abnormal wound healing, burns, infertility or diabetic shock. This polynucleotide expression of ADAMTS-SI in chondrocytes derived from osteoarthritic (OA) cartilage and induction by proinfilammatory cytokines.

Sequence 23 BP; 7 A; 7 C; 5 G; 4 T; 0 other;

Query Match Best Local S Matches 15 3 TGCGTTTGCCACTCAGAG Similarity Conservative 66.0**%**; 20 0 Score 13.2; DB 24; Pred. No. 2.4e+03; Mismatches 2; L-2.4e+03; 3; Length Indels 23; °, Gaps 0

N

RESULT 9 ABI83724/c

ABI83724 standard; DNA; 24 ВÞ

ABI83724;

15-FEB-2002 (first entry)

Capture oligonucleotide Zip ID#664 oligo #1.

Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious diseas infection; 21 hydroxylase deficiency; Turner Syndrome; obbesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss. disease;

Synthetic

WO200179548-A2

04-APR-2001; 2001WO-US10958.

14-APR-2000; 2000US-197271P.

(CORR) CORNELL RES FOUND INC

Barany F, Zirvi M, Gerry NP, Favis R, Kliman R;

Designing capture oligonucleotide probes for use on a support complementary oligonucleotides hybridize with little mismatch

Fig 25; 300pp; English

C(I) have melting temperatures within a narrow range. The method is useful CC (I) have melting temperatures within a narrow range. The method is useful CC e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus neoformans, Candida albicans and CC Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and pollo virus, and parasitic infectious agents CC selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects.

CC Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC p33 gene, human papillomavirus types 16 and 18 and liver cancers. The CC method is also used for environmental monitoring, forensics and the food cand feed industry, detecting comprises scanning (using e.g. a scanning CC electron microscope and infrared microscope) the support at the comparative and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to of the present invention. The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (II) have melting temperatures within a narrow range. The method is useful of the present invention.

Sequence 24 BP; 3 A; 9 C; 6 G; 6 T; 0 other;

Query Match Best Local Matches Similarity Conservative 64.0%; 0 Score 12.8; DB 24; Pred. No. 3.9e+03; 0; Mismatches 2; Length Indels .. Gaps 0

RESULT 10 ABI83725

밁

16

GGGATTGCCACTCAGA 1

S

4 GCGTTTGCCACTCAGA 19

ABI83725 standard; DNA; 24 BP

ABI83725;

15-FEB-2002 (first entry)

Capture oligonucleotide Zip ID#664 oligo #2.

Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.

Synthetic

WO200179548-A2

04-APR-2001; 2001WO-US10958

14-APR-2000; 2000US-197271P.

(CORR) CORNELL RES FOUND INC

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REPERENCE
AUTHORS
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AUTHORS
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BH854200
                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)

1 (bases 1 to 20)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A., and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                              AZ659612
20 bp DNA linear GSS 14
1M0537A07F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0537A07 F, DNA sequence.
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Class: TDNA tagged.
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Unpublished (2001)
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosida II; Brassicales; Brassicaceae; Arabidopsis.
plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Salk Institute Genomic Analysis Laboratory (SIGnAL) The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Joseph R. Ecker
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Fax: 858 558 6379
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                                                          Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10010 N. Torrey Pines Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lonso,J.M., Leisse,T.J., Barajas,P., Chen,H.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H.,
Zimmerman,J. and Ecker,J.R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html" 5 c 5 g 10 t
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/clone lib="Arabidopsis thaliana TDNA insertion lines"
/clone PPCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
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/strain="Columbia 0"
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Prednis, L., Shinn, I
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plasmid inserts
Unpublished (2000)
Contact: Robert B.
                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 21)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,B., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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Insert Length: 10000 Std
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                                                                                                                                                                                                                                                                                                                                         AZ663981
                                                                                                                                                                                                                                                                                                                                                         1M0543J22R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0543J22 R, DNA sequence.
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Fax: 801 585 7177
                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
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lgh quality sequence stop: 20.
Location/Qualifiers
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USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
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clone="UUGC1M0537A07"
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70.6%;
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RESULT 13
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human, mouse, alpha-catenin, primer, ss, cytostatic, antiinfertility, cadherin-catenin related pathway; heart testis, cancer; gene therapy; cadherin-catenin related disease; specifically dilated cardiomyopathy, cardiomyopathy; male infertility; CTNNA3; PCR; alpha T-catenin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      where the diagnostic method is an immunological method such as immunoblotting, serological tests or EIISA (enzyme linked immunoabsorbant assay). LppQ DNA and protein are useful for the preparation of a vaccine against infection by M. mycoides subsp. mycoides SC. LppQ DNA is useful for the preparation of live, attenuated or inactivated cells of M. mycoides subsp. mycoides SC to be used as marker vaccine, where the cells lack the ability to synthesise the LppQ lipoprotein. LppQ DNA is also useful for expressing LppQ protein in a detection method for M. mycoides subsp. mycoides SC, where the detection method is selected from PCR and hybridisation. LppQ protein is useful for detecting antibodies directed to it in a body fluid such as serun or bronchial fluid from an animal. LppQ protein are also useful as a specific antigen for sero-detection of contagious bovine pleuropneumonia (CBPP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a PCR primer used for site directed mutagenesis (to exchange Mycoplamsa specific UGARTTP codons to universal UGGTP codons) of lipoprotein LppQ gene from Mycoplasma mycoides subsp. mycoides SC (small colony type). LppQ is useful in a diagnostic method for the direct or indirect detection of M. mycoides subsp. mycoides SC, for the direct or indirect detection of M. mycoides subsp. mycoides SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel protein LppQ of Mycoplasma mycoides subsp. mycoides SC, parts useful for detection of Mycoplasma and for preparation for treating infection caused by Mycoplasma
                                                                                                                                                                                                                                                                                                                                                                             12-JUL-2000; 2000EP-0202472.
14-JUL-2000; 2000US-218309P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUN-2001; 2001WO-EP07392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 25
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                                                 New alpha catenin polypeptides and polynucleotides encoding the useful for predicting, diagnosing or treating cadherin-catenin diseases, particularly cardiomyopathies, cancer and male infer
   Examples;
                                                                                                                                                                                                                                                                                                              (VLAA-) VLAAMS INTERUNIVERSITAIR INST BIOTECHNOG.
                                                                                                                                                                                           2002-171717/22
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   Page
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                                                                                                                                                                                                                                                   Goossens S,
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36; 132pp;
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87.5%;
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Pred. No. 3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                   Vanpoucke G;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                  Diagnosing a subject at the risk for or having Alzheimer's disease comprises determining at least one single nucleotide polymorphism in mitochondrial DNA associated with the disease in the sample from the
                                                                                                                                                                                                                                                                                                                                                          subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Herrnstadt C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's disease-linked mitochondrial SNP PCR primer #262.
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99US-0160901.
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mtDNA; I
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Pred. No. 4.8e+03;
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The present invention describes a novel method for determining the risk of or disgnosing Alzheimer's disease using single nucleotide polymorphisms (SNPs) present in an individual's mitochondrial DNA (mtDNA). In additon, the SNPs identified can be used to identify agents suitable for use in treating Alzheimer's disease. Sequences AAC67301-C67610 are PCR primers used to demonstrate the method of the

Example 9;

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AZ339811/c
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Matches 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE
                                                                                                                                                                                                                                                                                                                                          BOUTCE
                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
Fax: 801 585 7177
Email: 6ddunægenetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0071 row: M column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     uunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ339811
AZ339811.1 GI:10414445
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 23.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso:
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0071M19"
(http://www.jax.org/resources/documencs/dnares/). The DNP was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                        clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                 from the Jackson
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ORIGIN
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TA106C01Q/c
                                                                                      BASE COUNT
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Query Match
Best Local S
Matches 10
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                                                                                                                                                                                                                      source
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T. brucei sheared genomic genomic survey sequence.
                                                                                                                                                                                                                                                                                                                             to give a tight size distribution (
4 kb). The v + 1 method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, exford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (10-DEC-2000) Trypanosoma brucel genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 ISA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trypanosoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AL461896.1
GSS.
                                                                                                                                                                                                                                                           Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma brucei.
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  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E: coli XL10-Gold (Stratagene) cells and selected for ampicilin resistance."
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                                                                                                                                /organism="Trypanosoma
/strain="TREU927"
/db_xref="taxon:5691"
/clone="106c01"
                                                                                                                                                                                                                                           Location/Qualifiers
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  0;
  Score 8.8; DB 17;
Pred. No. 1.6e+06;
0; Mismatches 2;
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Pred. No. 1.6e+06;
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ABK17746/c
ID ABK17746 standard; RNA; 17
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The invention relates to a nucleic acid molecule (I) which down regulates expression of an Ets-related gene (ERG). (I) is useful for treating conditions selected from cancer, lymphoma, Ewing's sarcoma, melanoma, tumour angiogenesis, diabetic retinopathy, macular degeneration, neovascular glaucoma, myopic degeneration, arthritis, psoriasis, verruca vulgaris, angiofibroma of tuberous sclerosis, port-vine stains, Sturge weber syndrome, Kippel-Trenaunay-Weber syndrome, Osler-Meber-rendu syndrome, leukaemia, osteoporosis and wound healing. (I) is useful for treating a patient having a condition associated with the level of ERG, by contacting cells of the patient with (I) under conditions suitable for the treatment. The method comprises the use of one or more therapies under conditions suitable for the treatment. Leukaemia or tumour anglogenesis is treated by administering (I) to the patient in confunction with one or more of other therapies such as radiation or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; hammerhead ribozyme; cytostatic; antitumour; antidiabetic; ophthalmological; antiarthitic; antipsoriatic; virucide; osteopathic; ophthalmological; antiarthitic; antipsoriatic; virucide; osteopathic; vulnerary; cancer; lymphoma; Eming's Barcoma; melanoma; psoriatei; tumour angiogenesis; diabetic retinopathy; macular degeneration; neovascular glaucoma; myopic degeneration; arthritis; verruca vulgaris; angiofibroma of tuberous sclerosis; port-wine stain; wound healing; sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; leukaemia; ss; Osler-Weber-rendu syndrome, leukaemia; osteoporosis; DNAzyme; inozyme;
                                                                                                                                                                                                                                                                       Claim 4; Page 65; 149pp; English.
                                                                                                                                                                                                                                                                                                                       Novel polynucleotide which down regulates expression of Ets-related gene, useful for treating cancer, diabetic retinopathy, macular degeneration, arthritis, psoriasis, verruca vulgaris and Sturge Web.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-082995/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-)
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Pred. No. 7.
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RESULT 18
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                                                                                                        AAQ90137 and AAQ90138 are a pair of primers for the PCR amplification of the transthyretin (TTR) gene exon 2. They were used in a PCR-SSCP analysis for the identification of the TTR mutation Gly6 to Ser6. This mutation, as a genetic marker for familial Alzheimer's disease, is useful for identifying people at risk of developing
                                                              Sequence
                                                                                                                                                                                                                                        Methods of useful for
                                                                                                                                                                                                                                                                                       WPI; 1995-185876/24.
                                                                                                                                                                                                                                                                                                                   Eisenberg-grunberg
                                                                                                                                                                                                                                                                                                                                                                                                                  03-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
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                                                                                            amyloidosis.
                                                                                                                                                                                                       Example 5; Page 32; 54pp;
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                Local
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14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23 bp DNA linear GSS 16-FEB-201
2M0018C03F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0018C03 F, DNA sequence.
                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duvai,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                   Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts Unpublished (2000) Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 23)
University of Utah Genome Center University of Utah
                                                                                                                                                                and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
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/note="Organ: lung; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using Zap-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
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/clone_lib="NIH_MGC_7"
/tissue_type="small_cell_carcinoma"
/cell_line="MGC3"
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/db_xref="taxon:9606"
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Hillier, L. Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Travaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 25)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA034350 ARNA linear EST 10-MAY-1 zk20301:r1 Soares pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:471120 5' similar to PIR.A44125 A44125 high density lipoprotein-binding protein, 110K - human ;, mRNA sequence.
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0018 row: C column: 03
                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                         AA034350.1 GI:1506159
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agazose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gil 4732114 [gb | AF129072:1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amplicillin resistance."
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Pred. No. 2.1e+06;
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27-OCT-1995;
27-OCT-1995;
27-MAR-1996;
   Use of factor
                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detecting and quantifying fungi and bacteria, involves obtaining a sequence of the fungus, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction and fluorescent probe
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                                                                                       (AMRA-)
                                                                                                                                                                                                                                                                                       eukaemia
                                                                                                                                                                                                                                                   sapiens.
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cytokine(s), especially oncostatin M and leukaemia inhibitory or functional derivatives or agonists - for the treatment or
                                                               ပ္ပိ
                                                                                                                                                                                                                                                                                                     cancer;
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                                                                                       AMRAD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page
                                                                                                                                                                                                                                                                                    icer; PCR primer; cytokine; oncostatin M; prophylaxis;
inhibitory factor; agonist; cell morphology; ss.
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                                                             Douglas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                       OPERATIONS
                                                                                                                                                                                                                                                                                                                          tissue cDNA PCR primer from the IL-3R alpha gene.
                                                                                                             95AU-0006242.
95AU-0006243.
96AU-0008966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14; 110pp; English.
                                                                                                                                                 96AU-0000364
                                                                                                                                                                           96WO-AU00676
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RESULT 23
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Best Local
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                       The present sequence invention describes ribozymes targeted to modulate the synthesis and/or expression of interleukin (II)-2R gamma encoded RNA. AAV93889 to AAV94574 represent specifically claimed ribozymes, and AAV94575 to AAV94574 represent specifically claimed substrate sequences from the present invention. The ribozymes can be used for the treatment of, e.g. graft rejection, autoimmune disease, cancer, psoriasis, allergy and other inflammatory conditions. The ribozymes are also used to induce tolerance in a recipient to alloantigen from a donor.
                                                                                                                                                                                            Ribozymes targeted to interleukin 2 - us cancer, autoimmune disease and allergies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; IL-2 receptor g-chain; interleukin 2 receptor gamma chain; hammerhead ribozyme; hairpin ribozyme; substrate; expression; can autoimmune disease; psoriasis; allergy; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A novel method has been developed for the treatment or prophylaxis breast cancer in a mammal. The method involves administering one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          prophylaxis of breast cancer by ameliorating delaying the onset of cancer.
                                                                                                                                                                Claim 4;
                                                                                                                                                                                                                                        WPI; 1998-333332/29
                                                                                                                                                                                                                                                                       McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                    03-DEC-1996;
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                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC.
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14; Conservative
                                                                                                                                                              Page 45; 61pp; English.
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Sequence 17 BP; 3 A;

7 C;

2 G;

5 U; 0 other;

8

TGCCACCGCGGGAGC

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RESULT 27
AA991484/c
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VERSION
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TA307A07Q/c
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                23
                                                                                                                                                                                                                                                   os91h03.sl NCI CGAP GC3 Homo sapiens cDNA clone IMAGE:1612757 3' similar to SW:FRPM HUMAN P10161 SALIVARY PROLINE-RICH PROTEIN PO CONTAINS MERZ2.tl MSR1 repetitive element ;, mRNA sequence. AA991484
                                                    Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini, Hominidae, Homo.

1 (bases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making smal insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shea
                   Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                  AA991484.1 GI:3177973
                                                                                                                                                                                                                                                                                                                                       AA991484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton Cambridge CB10 15A, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trypanosoma brucei
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Contact: Robert Strausberg,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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/strain="TREU927"
/db_xrefe="taxon:5691"
/clone="307a07"
a 11 c 4 g
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Pred. No. 2:1e+06;
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                                                                      CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.B. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

NCI-CGAP institute, Cancer Genome Anatomy Project (CGAP);
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Trace considered overall poor quality Seq primer: -40UP from Gibco
                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST.
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similar to SW:MGN_DROME P49028 MAGO NASHI
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                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTTCGGCGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; lst strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Ec RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is not normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone="IMAGE:1612757"
/clone_11b="NCI CGAP_GC3"
/tlssue_type="pooled_germ cell tumors"
/lab_host="DH108"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 8.4; DB 9;
Pred. No. 2.6e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 bp mRNA
omo sapiens cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     clone IMAGE:1947009 3' PROTEIN. ; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ph.D.,
                                                                                                                                                                                                                                             Ph.D., Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST 30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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The invention relates to a nucleic acid molecule which down regulates compression of a CD20 gene and a nucleic acid molecule which down compression of a neurite growth inhibitor gene (NOGO).

CC The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a CD nazyme) an Inozyme (an endolytic nucleic acids (e.g. a ribozyme or a CD nazyme) an Inozyme (cleaving RNA with an NGN triplet), a zinzyme (cleaving RNA with an Angulation and inflammatory arthropathy. The CD20 targetting concleic acid is used to cleave RNA of the NGCO gene in the presence of a cliving and inflammatory arthropathy. The NGCO-targetting concleic acid is used to cleave RNA of the NGCO gene in the presence of a cliving and inflammatory arthropathy. The NGCO-targetting concleic acid is used to cleave RNA of the NGCO gene in the presence of a cliving a patient having a condition associated with the level of NGCO. The treat a patient having a condition associated with the level of NGCO. The treatment may further comprise the use of one or more therapies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNAzyme; inozyme; G-cleaver; amberzyme; zinzyme; lymphoma; leukaemia; B-cell lymphoma; non-Hodgkin's lymphoma; HHI; lymphocytic leukaemia; human immunodeficiency virus; HIV associated NHI; mantie-cell lymphoma; MCL; immunocytoma; IMC; immune thrombocytopaenia; stroke; dementia; inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; muitiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; Parkinson's disease; ataxia; Huntington's disease; creutzfeldt-Jakob disease; muscular dystrophy; neurodegenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ss; antisense therapy; cytostatic; antiinflammatory; haemostatic; cerebroprotective; nootropic; neuroprotective; antiparkinsonian; muscular; CD20; neurite growth inhibitor gene; NOGO; hammerhead ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurite growth inhibitor gene useful for treating, e.g., lymphoma, leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human NOGO Zinzyme #322
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(BLAT/)
(MCSW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-607195/69
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28-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHOW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      central nervous system injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BLATT L.
MCSWIGGEN J.
CHOWRIRA B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 2000US-181797P.
; 2000US-185516P.
; 2000US-187128P.
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Query Match Best Local

Similarity

59.0%; 86.7%; 4 C; 7

Score 11.8; D Pred. No. 1.2e 0; Mismatches

.2e+04 DB 19;

Length Indels

20,

0

Gaps

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Conservative

0

Sequence

20

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G; 1 T; 0

other;

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RESULT 27
AAV23720/c
ID AAV237
XX AAV237
XX Lipase
XX reduce
XX reduce
XX Cleani
XX WO9808
XX WPI;
XX Uaria
PT Subst.
PT Perfo
XX Varia
PT Subst.
PT Perfo
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                                              PCR primers AAV23717-32 were used during the course of the invention. The specification describes Pseudomonas pseudoalcaligenes lipase variants. The variant lipases described in the specification has have better wash performance than the original lipase. It has improved removal of lipid stains, reduced calcium dependence, better compatibility with detergents or their components, increased hydrophobicity, altered substrate specificity and better one-cycle wash efficiency. The variant lipase is used in detergent and cleaning compositions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               In particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CNA stroke). Alzheimer's disease, dementia, multiple selerosis (NS), chemotherapy-induced neuropathy, amyotrophic lateral sclerosis (ALS), parkinson's disease, ataxia, Huntington's disease, Creutzfeldt-Jakob disease, muscular dystrophy, and/or other neurodegenerative disease states which respond to the modulation of NOGO expression. The present sequence is a zinzyme molecule of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lipase; variant; improved wash performance; removal; lipid stain; reduced calcium dependence; one-cycle wash efficiency; detergent; cleaning composition; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                         Variants of lipase from Pseudomonas containing specific substitutions - deletions or additions, having improved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Okkels JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-AUG-1996;
27-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV23720 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 BP; 3 A; 6 C; 3 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                            Disclosure; Page 74; 104pp; English
                                                                                                                                                                                                                                                                                                                                                                     performance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NOVO ) NOVO-NORDISK AS
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96DK-0000902
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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SOURCE
ORGANISM
                                                                                                                                    REFERENCE
AUTHORS
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VERSION
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TA219C09P
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                                                                                                                                                                                                                                                                                                                                                   DEFINITION
                                                                                                                                                                                                                                                               KEYWORDS
                                                                                                                                                                                                                                                                                                                                                                         Spoo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 9; Conserv
                                                                      TITLE
                                                  JOURNAL
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                                                                                                                                                                                                                                                           AL479371 GI:11845122 GSS.
Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G. Direct Submission Submitted (10-DEC-2000) Trypanosoma brucel genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    æ
                                                                                                                                                                                         Trypanosoma brucei.
Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: |
Plate: 0294 row: P column: 19
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                           genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                               T. brucei sheared genomic
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Fax: 801 585 7177
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                                                                                                                                                                           Trypanosoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CGCTGGGAAC 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWP42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. Coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC1M0294P19"
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Pred. No. 2.6e+06;
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1M0294A18F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0294A18 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 TIGCTICGGC 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.or.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to give a tight size distribution ( 4\ kb ). The v + i method used for the library construction described in detail in Smith, H. and Venter, J.C. (Making (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nhl@Banger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0294 row: A column: 18
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center 
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B.
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igh quality sequence stop: 20.
Location/Qualifiers
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/strain="TREU927"
/db_xref="taxon:5691"
/clone="219c09"
a 8 c 4 g
                                                                                                                                                                                                                                                                                                                                                                                          Biomedical
                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0294A18"
                    /clone lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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'lab_host="E. Coli strain XL10-Gold, T1-resistant,
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18-MAY-1999;
13-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a recombinant DNA construct of a plant (Arabidopsis thaliana) centromers. The constructs are useful for producing stably inherited michrosomes which can serve as vectors for the construction of transgenic plant and animal cells expressing selected proteins such as hormones, enzymes, interleukins, clotting factors, cytokines, antibodies, and growth factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant DNA construct comprising a plant centromere, producing stably inherited michrosomes which can serve at the construction of transgenic plant and animal cells -
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                                                                                                                                                                                                                                                                                                                                                                                             JP2001321190-A
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Claim 4; Page 43; 528pp; Japanese
                                                Arraying genome clones
                                                                                                                                                                                                                                                                                   12-MAR-2001; 2001JP-0068285.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Keith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 11.8; DB 21;
Pred. No. 1.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for
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RESULT 32
AAH24199/c
ID AAH241
XX AAH241
XX AAH241
XX PCR pr
XX MOUSE,
KW FICOMB
KW Iigand
KW PCR pr
XX WO2001
XX Synthe
XX I) -NOV
XX II -NOV

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates (b) a primer designed based on the chromosome marker completel plates; (b) a primer designed based on the chromosome marker completel plates corresponding to the marker is detected from the resultant complified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified discrimination Nos. to array the multiwell complates; (e) the clones are mixed respectively in each wells of longitudinal cand lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell complates are specified from the detected result; and (i) the clones are mixed products and staged. The mixed products are the positions on the chromosome and arrayed. The nitroarray is useful for gene analysis. ABI42357 to ABI46332 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                               The invention relates to 3 novel mouse proteins which inhibit ageing (AAB73664-AAB73666) and to the nucleic acids encoding them (AAH24193-AAH24195). The invention also relates to recombinant vectors, host
                                                                                                                                                                                                                                                                                                                                                                        Shiraishi N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse; ageing inhibition-related DNA; ageing inhibitory protein; recombinant production; transgenic animal; detection; drug screening; ligand identification; ageing-related disorder; premature ageing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH24199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH24199 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primers for human chromosome 1p36-35 DNA, and ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention.
                                                                                                                                                                    Example 1; Page 89; 94pp; Japanese
                                                                                                                                                                                                                                       Polypeptide for treatment of premature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-NOV-2000; 2000WO-JP08121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200138529-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                           (KYOW ) KYOWA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes a method of arraying genome clones. hod comprises: (a) clones of the genomic libraries contained in tivell plates numbered for discrimination are mixed in each of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTTTGCCACTCAGAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       накко косуо кк
                                                                                                                                                                                                                                                                                                                                                                     Sekine S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99JP-0329649
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                                                                                                                                                                                                                                                                                                                                                                        Nabeshima
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Pred. No. 1.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to isolate ageing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                     Fujimori
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                                                                                                                                                                                                                                                                                                                                                                        'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inhibition-related DNA.
   DNA encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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cells and

animals

comprising

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RESULT 36
AZ355490
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VERSION
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ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JOURNAL
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                                                                                                                                                                                          воитсе
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0095 row: N column: 14
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1M0095N14F Mouse 10kb plasmid UUGC1M clone UUGC1M0095N14 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 TIGCTICGGCGGG 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia, Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ355490.1 GI:10468057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 48.2%;
Similarity 76.9%;
10; Conservative
                                                                                                                                                                                                                                                                     .ass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGCTITGGTGTG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      308,
                                                                                                                                                                                                              quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi[4732114]gb[AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-completent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                        /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGCIM0095N14"
'lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                              sex="Male"
                                                  clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 8.2; DB 17;
Pred. No. 3.3e+06;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                              Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
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KEYWORDS SOURCE

밁 ફ

REFERENCE

AUTHORS

COMMENT

TITLE

FEATURES

Query Match
Best Local Similarity 76.
Matches 10; Conservative

48.2%; Score 8.2; DB 9; 76.9%; Pred. No. 3.3e+06; tive 0; Mismatches 3

Length 22;

Indels

0

Gaps

0

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SOURCE
ORGANISM
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VERSION
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AA908627
                                                                                                                                                                                                                                                                                                                           FEATURES
  BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMMENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUTHORS
                                                                                                                                                                                                                                                                                                  source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
Eukaryota; Metrazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                                                                                                                                                                                                           unknown library type
Trace considered overall poor quality
Seq primer: -40m13 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       og86e01.s1 NCI_CGAP_Ov8 Homo sapiens cDNA clone IMAGE:1455192 3 similar to SW:ULA6_HUMAN P30041 24 KD PROTEIN ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 1,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA908627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA908627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA908627.1 GI:3048032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTTGCTTCTATGG 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 Kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                               /note="Organ: ovary; Vector: pCMV-SPORT4; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.1 kb. Life Technologies catalog #:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/dlone="IMAGE:1455192"
/clone="IMAGE:1455192"
/clone_Tib="NCI_CGAP_Ov8"
/clssue_type="serous_adenocarcinoma"
/tissue_type="serous_adenocarcinoma"
/tab_host="DH10B"
                              10982-015"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     48.2%;
76.9%;
     <u>4.</u>
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Pred. No. 3.3e+06;
0; Mismatches 3;
7
9
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RESULT 35
ABI88869
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                   CC oligonuclectide probes (I) for use on a support to which complementary CC oligonuclectide probes (II) will hybridise with little mismatch, where CI have melting temperatures within a narrow range. The method is useful CC for detecting infectious diseases caused by bacterial infectious agents e.g. Cryptococcus necformans, Candida albicans and CC infectious agents e.g. Cryptococcus necformans, Candida albicans and CC Aspergillus fundgautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and polio virus, and parasitic infectious agents CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. CC etecting cancer involving oncogenes, tumour suppressor genes, or genes cinvolved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the CC sets occurred and correlating (using if lyation of the oligonucleotide probe genes or absence of the target nucleotide sequences. ABIS2074 to affect in present inventor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious diseas infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABI88869
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                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Fig 25; 300pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-034366/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Barany
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            the present
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LSJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zirvi M,
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86.7%;
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Pred. No. 1.3e+04;
0; Mismatches 2;
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RESULT 36
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AC AAQ342
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                                                                                                                      The sequence shows an upstream PCR primer for in vitro amplification of bovine microsatellite sequences obtd. by screening library of bovine MboI DNA fragments of between 250 and 500 by with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-thybridsed. Assuming independent distribution of microsatellites and MboI sites, the frequency of (TS)n > microsatellites in the bovine genome is estimated at >100,000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). For each such microsatellite sequence sufficient information was obtd. to microsatellites microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding.
                                                Best Loc
Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                       gene mapping, and selective breeding
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                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-284684/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Georges M, Massey JM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-AUG-1992
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                                                                                                                                                                                                                                                                                                                                                             Table 8; Page 413; 517pp; English
                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JAN-1992;
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19
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13; Conserv
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                                                             Similarity
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TTTGCCTGTGCCAAGCAG 2
                      TTTGCGTTTGCCACTCAG 18
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                                                                                                  B₽;
                                                Conservative
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86.7%;
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                                                            58.0%;
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                                                            Score 11.6; DB 13;
Pred. No. 1.6e+04;
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Pred. No. 1.3e+04;
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                                                Mismatches
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                                                                                                   other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 24;
                                                                       Length 19;
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JOURNAL COMMENT
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AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOURCE
ORGANISM
Search completed: June 7, 2003, 09:43:43 Job time: 1020.33 secs
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                                                                                                                                                                                                                                                                                                                                    BASE COUNT
ORIGIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PEATURES
                                                                                                                                                                                                                                              Query Match 48.2%;
Best Local Similarity 76.9%;
                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606

Pax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00

Plate: 0028 row: H column: 03

Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                     10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 23)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A., Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                         4 GCTTCGGCGGGAA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 house mouse.
Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 23.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmid inserts
                                                                                                         GCTGAGGGGGGAA 22
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwh92 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coil XL10-Gold (Stratagene) cells and selected for ampicillin resignance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ASU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0028H03"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       aboratory Mouse DNA Resource
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                                                                                                                                                                                                               Score 8.2; DB 17;
Pred. No. 3.3e+06;
0; Mismatches 3;
                                                                                                                                                                                                                                                                  Length 23;
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RESULT 40
ABL43246
ID ABL43
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Best Local Similarity
Matches 14, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anti-immunodeficiency virus immunokine (a neurotoxin) capable of binding to a cellular protein in a manner that prevents HIV infection of the cell. One such neurotoxin is alpha-cobratoxin from Naja naja siamensis (Thailand cobra). The present sequence is a PCR primer which was used to amplify alpha-cobratoxin cDNA for use in the present invention.
                                                                                                                                                                                                                                                           Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Composition for preventing HIV infection of mammalian cells comprises an anti-immunodeficiency virus immunokine capable of binding to a cellular protein by which HIV infection of the cell is prevented -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-616350/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mundschenk DD, Reid PF;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-MAR-2001; 2001WO-US08150.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200170173-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Naja naja siamensis
                                                                                                                                                                                                        JP2001321190-A.
                                                                                                                                                                                                                                                                                                   Human chromosome 1p36-35 PCR primer SEQ ID NO:290.
                                                                                                                                                                                                                                                                                                                                11-APR-2002
                                                                                                                                                                                                                                                                                                                                                           ABL43246;
                                                                                                                                                                                                                                                                                                                                                                                     ABL43246 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 4 A; 6 C; 3 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a composition for preventing HIV infection of mammalian cells. The composition comprises an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 30; 54pp; English.
                                                                                                                                                                             20-NOV-2001.
                                                                                                                                                                                                                                  Homo sapiens
Claim 4; Page 10; 528pp; Japanese.
                          Arraying genome clones
                                                                                                                        10-MAR-2000; 2000JP-0066716
                                                                                                                                                 12-MAR-2001; 2001JP-0068285
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                                                     WPI; 2002-144136/19.
                                                                                (RIKA ) RIKAGAKU KENKYUSHO (GENO-) GENOTEX YG.
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The present invention describes a method of arraying genome clones. The CC multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (a) a primer designed based on the chromosome marker contained in the clones is added to the mixture to carry out an amplification resultant confided product to specify the discrimination Nos. of the multiwell confided product to specify the discrimination Nos. of the multiwell confided product to specify the discrimination Nos. of the multiwell confided plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell confided; (e) the clones in the multiwell plates of the specified confided and lateral directions; (f) the maximum the absorbance of longitudinal and lateral directions; (f) the mixed clones are cultured and the created from the amplified by using the above primer; (g) signals created result; and (i) the clones are constituted as the positions on the chromosome and arrayed. The constituted as the positions on the chromosome and arrayed. The creonestituted as the positions on the chromosome and arrayed. The creonestituted as the positions on the chromosome and arrayed. The constitutes for human chromosome 1936-35 NNA, and ABL45322 represent pCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention.

XX Sequence 21 BP; 3 A; 9 C; 1 G; 8 T; 0 other;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 25)
               University of Chicago Medical Center 5841 S. Maryland Ave., MC2115, Chicago, Tel: 773-702-6788
                                                                                          Zhou, G., Chen, J., Lee, S., Terry, C., Rowl
The pattern of gene expression in human
stem/progenitor cells
Unpublished (2001)
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Direct Submission

Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CBI0 1SA, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk and constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucesi (TREU927/4 GUTat 10.1) was mechanically sheared to give a tight size distribution (

4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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This EST fragment was amplified from cDNA Library of human CD 34+
stem/progenitor cells with GLGI technique (Generation of Longer
CDNA fragments from SAGE tags for Gene Identification, Proc. Natl.
Acad. Sci. USA 97, 349, 2000), which starts from the 3' end till
the first CATG site of the target cDNA sequence.
Seq primer: Ml3 Forward.
                                                                                                                                                Email: neleayed@tigr.org
Details of T. brucei secuk/Projects/T_brucei/.
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                              Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Trypanosoma brucei
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                                                         /organism="Trypanosoma b
/strain="TREU927"
/db_xref="taxon:5691"
/clone="352b06"
a 4 c 13 g
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/db_xref="taxon:9606"
/clone_lib="cDNA library of human CD 34+ stem/progenitor
cells"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="Bone marrow"
/cell_type="CD34+ stem/progenitor cells"
/note="3'ES9s converted from the SAGE tag sequences using
GLGI method"
4 c 5 g 13 t
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 Score 10.4; DB 17;
Pred. No. 2.4e+05;
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Pred. No. 9.1e+04;
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Best Local S
Matches 15
                                                                                                                                                                                                                                                  comprises deter DNA variant, eq disequilibrium
       The present invention describes a method (M1) of determining an individual's predisposition to epilepsy and/or development of epilepsy, as well as predicting the individual's response to medication. The method comprises determining the genotype of at least one gene selected from SCNIA, SCNZA or SCN3A, or a DNA variant, equivalent, or mutation which shows a linkage disequilibrium. SCNIA, SCNZA and SCN3A are all sedium channel genes located on chromosome 2. The idiopathic generalised epilepsy (IGE) gene is more specifically localised on chromosome 2q23-q31. Compounds identified as modulators of the biological activity of SCN1A, SCNZA or SCN3A proteins or genes, are useful for treating epilepsy or other neurological disorders. They have anticonvulsant and
                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of detecting and quantifying fungi and bacteria, involving obtaining a sequence of the microorganism to be detected and quantified, extracting the DNA from the sample, and subjecting the DNA to polymerase chain reaction (PCR) and fluorescent probe analysis. The method is useful for identifying and quantifying specific fungi and bacteria using specific DNA sequences. The specific DNA sequences are useful for the real time detection of PCR products with a fluorogenic probe system or other molecular probes like hybridisation. ABK29026-ABK29474 represent fungal and bacterial PCR primers and probes used in the method of the invention.
                                                                                                                                                                                                              Example 5; Fig
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                                                                                                                                                                                                                                                                                                                                                                      Rouleau GA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; epilepsy; chromosome 2; SCN1A; SCN2A; SCN3A; identificati diagnosis; mutation; chromosome 2q23-q31; neurological disorder; anticonvulsant; neuroprotective; PCR primer; ss.
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2.9e+02
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transetted to the human Nck-2 gene, which inhibit the expression.

CC The antisense oligonucleotides were designed to target different

CC regions of the human Nck-2 mRNA, and were analysed for their effect on

CC Nck-2 mRNA levels by quantitative real-time PCR. Nck-2 (also known

CC as Nck adapter protein, hNck-beta and Grb4), contains both SH2 and SH3

CC arc homology domains and functions as an adapter protein in

CC integrin-mediated and receptor tyrosine kinase-mediated signal

CC transduction, particularly in growth factor receptor signalling.

CC Moreover, Nck-2 participates in pathways that connect growth factor

CC receptor signalling and integrin signalling via its interaction with

CC integrin, growth factor and Wnt signalling val its interaction with

CC integrin, growth factor and Wnt signalling pathways. Nck-2 also

CC interacts with EGF (epidermal growth factor) and PDGF-stimulated DNA

CC growth factor) receptors, inhibiting EGF- and PDGF-stimulated DNA

CC v-Abl, Ras and Sos proteins to induce transcriptional activation, and is

CC therefore implicated in the development of cancer, particularly leukaemia

CC and breast cancer. The oligonucleotides of the invention are useful for
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Best Local !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antisense compound which inhibits expression of human nck-2 useful for treating disease or condition associated with expression nck-2, and as research reagents, kits and diagnostics -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequences AAC92649-C92728 represent antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Column 41-42; 38pp; English.
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83.3%;
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Pred. No. 5.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pirect Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
Submitted (10-DEC-2000) Trypanosoma brucei genome Campus, Hinxton,
project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
nhl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma brucei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger at http://www.sanger.ac.uk/Projects/T_brucei/.
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                                                                                                                    Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R. and Wright, D., Weise, R. Mouse, whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                         2M0032B10F Mouse 10kb plasmid \bar{U}UGC1M library Mus musculus clone \bar{U}UGC2M0032B10 F, DNA sequence.
                                                                                                                                                                                                                                                                                             Mus musculus
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                                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                   AZ786547.1
                 University of Utah Genome Center University of Utah
                                                                                Unpublished (2000)
                                                                                                  plasmid inserts
                                                                                                                                                                                                                                                                                                                        house mouse.
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/strain="TREU927"
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/clone="114g12"
5 c 7 g
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Polymers Research Bldg.,
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Pred. No. 6.1e+05;
0; Mismatches 4;
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Insert Length: 10000 Std Error:
Plate: 0032 row: B column: 10
Seg primer: CGTTGTAAAACGACGGCCAGT
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High quality sequence stop: 24.
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Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                     Tetrahymena thermophila.
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                  5009-0-4-D05.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GITGCTTCGGC 11
                                                                                                                             J. and Klobutcher,L.
EST from Terrahymena thermophila,
Unpublished (2002)
Contact: Turkewitz AP
                    Molecular Genetics and Cell Biology
University of Chicago
920 B. 58th Street, Chicago, IL 606
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                               1 (bases 1 to 18)
Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E.,
                                                                                                                                                                                                                                                                                                                                                                     BM398017.1 GI:18198070
                                                                                                                                                                                                                                                                                                                                                                                           BM398017
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwd42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
    apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
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/clone="UUGC2M0032B10"
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/strain="C57BL/6J"
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                                                                Chicago, IL 60637,
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Pred. No. 7.9e+05;
0; Mismatches 1;
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Length 24; Indels

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from the Jackson

linear

EST 17-JAN-2002

growing cells

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RESULT 7
AAV35581,
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McSwiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequences (AAQ50775-99) and (AAQ52551-8) are pref. hepatitis B virus target sequences for enzymatic Randlecules. The RNA molecules are complementary to a substrate binding region in the specified gene target. They also have enzymatic activity, in that they specifically cleave RNA in the target. The ERMS interfere with viral replication and therefore have anti-viral properties. They can be used to attenuate viruses to be used in vaccines.
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                                                                                                 Hydronephrosis gene - useful to treat or diagnose renal diseases disorders, e.g. vesical-ureteral reflux, pelvi-ureteral junction obstruction, multicystic renal dysplasia or renal agenesis
                                                                                                                                                                                                                                                                                                         WO9815650-A2
                                                                                                                                                                                                                                                                                                                                                                               vesical-ureteral reflux; pelvi-ureteral junction obstruction; multicystic renal dysplasia; renal agenesis; hydronephrosis; von Mayer-Rokitansky-Kuester disorder; bifid ureter; STS prob
                                       This primer probes used
                                                                                                                                                     WPI; 1998-240833/21
                                                                                                                                                                                                                              09-OCT-1996;
                                                                                                                                                                                                                                                       09-OCT-1997;
                                                                                                                                                                                                                                                                                16-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                       Hydronephrosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                 STS probe GV10
This primer is used for the generation of sequence tagged sites (STS) probes used for pinpointing chromosome 6 breakpoint. This was used fo isolating the human hydronephrosis (HNG) gene of the invention. A translocation partner to this HNG gene on chromosome 6 is the chromosome 19 USF2 gene. The HNG gene can be used as a starting point
                                                                            Example 3;
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                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                          Page 46; 73pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 4 A; 6 C; 4 G; 3 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                       site; PCR primer; ss.
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                                                                                                                                                                                                     BIOTECHNOG.
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RESULT 8
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ID AAK98779;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KW antiasthmatic; neuroprotective; immunomodulator; cytostatic; vasotropic;
KW antialergic; dermatological; antiarteriosclerotic; cerebroprotective;
KW antialergic; dermatological; antiarteriosclerotic; cerebroprotective;
KW cardiant; immunosuppressive; anticonvulsant; antiparkinsonian; analgesic;
KW antinigraine; antidepressant; ophthalmological; vulnerary; antidiabetic;
KW arthitis; inflammatory bowel disease; Crohn's disease; atherosclerosis;
KW periodical disease; emphysema; acute respiratory distress syndrome;
KW arthitis; inflammatory bowel disease; Alzheimer's disease;
KW organ transplant toxicity; rejection; cachexia; allergy; restenosis;
KW cancer; tissue ulcerations; epidermolysis bullosa; aortic aneurysm;
KW osteoporosis; artificial joint implant; congestive heart failure; stroke;
KW myocardial infarction; cerebral implant; congestive heart failure; stroke;
KW depression; partificial joint implant; congestive heart failure; stroke;
KW depression; peripheral neuropatha; head trauma; corneal injury;
KW hortopic; cognition enhancement; amyotrophic lateral sclerosis;
KW coular angiogenesis; macular degeneration; abnormal wound healing; burn;
KW mortogic; cognition enhancement; chondrocyte; osteoarthritic; OA; cartilage;
KW mridellammatory cytokine, human, corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to design suitable compounds or techniques for the treatment of renal diseases or disorders, or nucleotide probes for diagnosing cells involved in renal diseases or disorders. A protein or a fragment
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   Novel isolated polynucleotide encoding A Disintegrin And Metalloprotease family of metalloproteases, containing to domain, termed ADAMTS-SI, useful for diagnostic, biother
                                                                                                                                                                                                        Buckbinder L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense primer for amplification of ADAMTS-SI.
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                                                                                                                                                                                                            Mitchell PG,
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JOURNAL
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                                                                                                                                                                                                                                                                               Bource
                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                nhiseanger.ac.uk

Constructed at the Institute for Genomic Research (TIGR),
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Rypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects. In
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Trypanosoma brucei.
Trypanosoma brucei
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GSS.
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T. brucei sheared genomic
                                                                                                                                                                                                                                                                                              Email: neleayed@rigr.org
Details of T. brucei sequencing at the Sanger
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genomic survey sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
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GTTTCTTCGGTAGG 18
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                                                                                                                                                                            /clone="75a03"
4 c
                                                                                                                                                                                                                                        organism="Trypanosoma/
strain="TREU927"
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78.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    source
BH854200 25 bp DNA linear GSS 13-JUN-2002 SALK 078881.34.80.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK 078881.34.80.x, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
Contact: Marra M/WashU-NCI Mouse EST Project 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AI594892 25 bp mRNA linear EST 21-APR-1999 vel2c06.xl Soares mouse NbMH Mus musculus cDNA clone IMAGE:817930 37 similar to SW:SUR_RAT Q09429 SULFONYLUREA RECEPTOR. [1] ;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 25).
Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T.,
Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T.,
Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person
,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter
,B., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R.,
Waterston,R. and Wilson,R.
The WashU-NCI Mouse EST Project 1999
                                                                                                                                           J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314,286 1800

Pax: 314 286 1810

Email: mouseest@watson.wustl.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Trace considered overall poor quality possible reversed clone: similarity or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This clone was previously sequenced on the 5' data is from the 3^{\prime} end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MGI:486210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AI594892.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nouse mouse
                                                                                                                                                                                                                                       Similarity
                                                                                                                                       GCTTCTGCTGGTAC 20
                                                                                                                                                                              GCTTCGGCGGGAAC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop: 1.
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                           /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="IMAGE:817930"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tissue_type="heart"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone_lib="Soares mouse NbMH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GI:4603940
                                                                                                                                                                                                                                       54.1%;
78.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    stage="4 weeks"
host="DH108"
                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                       Score 9.2; DB 9; Pred. No. 1e+06;
                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             on wrong strand
                                                                                                                                                                                                                                                         Length 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         end only, this new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  contact
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                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch -
                                                                                                                                                                       misc_feature
                                                                                                                                                                                                                        Mycoplasma mycoides subsp. mycoides
                                                                                                                                                                                                                                                        Lipoprotein; LppQ; antibacterial; vaccine; Mycoplasma infection; bovine pleuropneumonia; CBPP; site directed mutagenesis; PCR primer;
                                                                                                                                                                                                                                                                                                         M. mycoides LppQ gene site directed mutagenesis PCR primer MMMLP482.
                                                                                                                                                                                                                                                                                                                                               06-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                 AAD07981,
                                                                                                                                                                                                                                                                                                                                                                                                                  AAD07981 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Fig 25; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-034366/04
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              24-NOV-2000; 2000WO-EP11798
                                                                                  WO200140471-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present invention describes a method (M1) for designing capture
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGGATTGCCACTCAGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GCGTTTGCCACTCAGA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 6 A; 6 C; 9 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
                                                                                                                                                                     Location/Qualifiers
                                                                                                              /*tag= a
/note= "Nucleotide G is present at this location in the
sequence shown in sequence listing of the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           64.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gerry NP, Favis R, Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.8; DB 24;
Pred. No. 3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                        SC.
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RESULT 12
AAD07982/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel protein LppQ of Mycoplasma mycoides subsp. mycoides SC, or its parts useful for detection of Mycoplasma and for preparation of vaccine for treating infection caused by Mycoplasma -
              WPI; 2001-381385/40
                                                                                                                                                                                                                                                              Mycoplasma mycoides subsp. mycoides SC
                                                                                                                                                                                                                                                                                                                                                      M. mycoides LppQ gene site directed mutagenesis PCR primer MMLP483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD07982 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 25 BP; 2 A; 5 C; 3 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 26; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-NOV-1999;
                                              Frey JJ,
                                                                                                                     29-NOV-1999;
                                                                                                                                                       24-NOV-2000; 2000WO-EP11798.
                                                                                                                                                                                             07-JUN-2001
                                                                                                                                                                                                                                WO200140471-A1.
                                                                                                                                                                                                                                                                                                  Lipoprotein; LppQ; antibacterial; vaccine; Mycoplasma infection; bovine pleuropneumonia; CBPP; site directed mutagenesis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD07982;
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                                                                                  (ALKU ) AKZO NOBEL NV
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                                              Nicolet JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TIGITITIGCCACTCA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                     99EP-0123676.
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87.5%;
                                                  Abdo EE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.8; DB 22;
Pred. No. 3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25;
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SOURCE
ORGANISM
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BM399936
                                                              TITLE
JOURNAL
COMMENT
                                                                                                                                                                          REFERENCE
                                                                                                                                                                                                                                                                                                                                                                      DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURES
                                                                                                                                                                                                                                                                                 KEYWORDS
                                                                                                                                                                                                                                                                                                                            ACCESSION
                                                                                                                                                                                                                                                                                                          /ERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                   AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12;
                                                                                                                                                                          EST.
Tetrahymena thermophila.
Tetrahymena thermophila
Tetrahymena thermophila
Tetrahymena thermophila
Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Eukaryota; Alveolata; Tetrahymenina;
                                                          1 (bases 1 to 25)
Turkewitz,A.P., Karrer,K.M., Jahn,C., Orias,E., Kirk,K.E., Frank
,J. and Klobutcher,L.
EST from Tetrahymena thermophila, strain CU423.1, growing cells
Unpublished (2002)
Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                 BM399936.1 GI:18199989
EST.
                                                                                                                                                                                                                                                                                                                            Tetrahymena thermophila cDNA, mRNA sequence. BM399936
                                                                                                                                                                                                                                                                                                                                                                5009-0-63-D08.t.1 Chilcoat/Turkewitz cDNA (large fraction)
                                                                                                                                                                                                                                                                                                                                                                                          ВМ399936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Insert Length: 10000 Std Error: 0.00 Plate: 0543 row: J column: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
                    Molecular Genetics and Cell Biology University of Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTTGCTTCGGCGGGAAC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGTGCTTCATGGGTAAC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was but end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt end-repaired and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWn92 (gil4732114|gb)AF129072.l), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
  58th Street,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PMP42nv; Purified genomic DNA from M. musculus C57BL/60 (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="UUGC1M0543J22"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52.9%;
70.6%;
Chicago,
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Pred.
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  IL 60637, USA
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                                                                                                                                                                                                                                                                                                                                                                                          mRNA
                                                                                                                                               Orias, E., Kirk, K.E., Frankel
                                                                                                                                                                                                                                                                                                                                                                                          linear
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 bp DNA linear GSS 04-OCT-200100298C06R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0298C06 R, DNA sequence.
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Fax: 773 702 3172
Email: apturkew@midway.uchicago.edu
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD$27N; Purified genomic DNA from M.
musculus C578L/62 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through o
0.005 inch orifice at constant velocity. The sheared DNA
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                                                                                                                                                                                                                                                                                                                                  /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0298C06"
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Pred. No. 1.3e+06;
0; Mismatches 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplify cDNA sequences encoding antigens of structural and non-
structural regions of non A non B hepatitis virus (NANBEY). Th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page
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                         12-MAR-2002
                                                                          ABK02611,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplified sequences were derived from the plasma of a NANBH patient by recombinant DNA techniques. The amplified fragments are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-1994.
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                                                                                                                     ABK02611 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences given in AAQ58828-72 are primers which were used to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosis of NANBH patients and
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SAKWA KAGAKU KENKYUSHO CO.
TONEN CORP.
ZH TOKYOTO RINSHO IGAKU SOGO KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                      Similarity
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Nucleic acid molecules, e.g., enzymatic nucleic acids and antisense constructs, which down regulate expression of a CD20 gene or neurity growth inhibitor gene useful for treating, e.g., lymphoma, leukemia
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                                                                                                                                                                                                                                                                                                                                                                                         Claim 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-607195/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-FEB-2000; 2000US-181797P
28-FEB-2000; 2000US-185516P
06-MAR-2000; 2000US-187128P
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammatory arthropathy; central nervous system injury; cerebrovascular accident; CVA; Alzheimer's disease; multiple sclerosis; chemotherapy-induced neuropathy; amyotrophic lateral sclerosis; ALS; parkinson's disease; ataxia; Huntington's disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Creutzfeldt-Jakob disease; muscular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BLATT
                                                                                                                                                                                                                                                                                                                                                                                        Page 137; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McSwiggen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chowrira BM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dystrophy; neurodegenerative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                leukemia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ribozyme;
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to cleave RNA of CD20 in the presence of a divalent cation that is preferably Mg^2+. Furthermore, it may be contacted with a cell to reduce CD20 activity of the cell and treat a patient having a condition associated with the level of CD20. The treatment may further comprise the used of one or more therapies. In particular, the CD20 targetting nucleic acid may be used to treat lymphoma, low-grade or follicular non-Hodgkin's lymphoma (NHL), bulky low-grade or follicular NHL, lymphocytic leukaemia, HIV (human immunocytoma (HVC), small B-cell lymphore, cell lymphoma (MCL), immunocytoma (HVC), and in sesciated NHL, mantle-cell lymphoma (MCL), immunocytoma (HVC), and in sesciated NHL, mantle-cell lymphocytom thrombocytopaenia, and inflammatory arthropathy. The NOGO-targetting nucleic acid is used to cleave RNA of the NOGO gene in the presence of a divalent cation that is preferably Mg^2+. Furthermore, the nucleic acid may be contacted with a cell to reduce NOGO activity of the cell and treat a patient having a condition associated with the level of NOGO. The treatment may further comprise the use of one or more therapies.

The particular, the NOGO-targetting nucleic acid may be used to treat central nervous system (CNS) injury and cerebrovascular accident (CVA, attrobe) altherer's disease demonstrate multithe sclear accident (CVA). The invention relates to a nucleic acid molecule which down regulates expression of a CD20 gene and a nucleic acid molecule which down regulates expression of a neurite growth inhibitor gene (NGGO). The nucleic acids may be enzymatic nucleic acids (e.g. a ribozyme or a DNAzyme) an Inozyme (an endolytic nucleic acid cleaving a an RNA molecule possessing an NCH motif), a G-cleaver (cleaving RNA with a NYN motif) pr an amberzyme (cleaving RNA with an NCN triplet), a zinzyme (cleaving RNA with an NCN triplet) a zinzyme (cleaving RNA with an NCN triplet) a zinzyme (cleaving RNA with an NCN triplet). stroke), Alzheimer's disease, de chemotherapy-induced neuropathy, Parkinson's disease, ataxia, Hun dementia, multiple sclerosis (MS) hy, amyotrophic lateral sclerosis al sclerosis (ALS) Creutzfeldt-Jakob

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SOURCE
ORGANISM
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                                                                                                 AUTHORS
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Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                       Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 20)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly

,M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.,

and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                  20 bp DNA linear GSS 29-SEP-20 1M0080L18F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0080L18 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTAt 10.1) was mechanically sheared to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L., Melville, S.B., Rajandream, M.A. and Barrell, B.G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hi Cambridge CB10 18A, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AL488313.1 GI:11864165
GSS.
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    whole genome scaffolding with paired end reads from 10kb
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/clone="266d03"
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Pred. No. 1.6e+06;
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                                                                     REFERENCE
AUTHORS
TITLE
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ORGANISM
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0080 row: L column: 18
Seq primer: CGTTGTARAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
Contact: Robert B
Unpublished (1999)
Other ESTs: 2820485.5prime
Contact: Robert Strausberg,
                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 22)
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Fax: 801 585 7177
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University of Utah
                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                      AW247820.1 .GI:6590813
                                                                                                                                                                                                                                                                                AW247820
                                                                                                                                                                                                                                                                                                   mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from musculus C57BL/6J (male) was obtained from the Ja Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                     22 bp mRNA linear EST 07-JAN-2000 Homo sapiens cDNA clone IMAGE:2820485 3',
         Ph.D
                                                                     Mammalian Gene Collection (MGC)
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RESULT 19
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Best Local S
Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Consensus; resistance; fungus; pathogen; wheat; cell wall; apposition; papilla; contact site; callose; carbohydrate; phenol; transgenic plant; Mlo; Erysiphe graminis; powdery mildew; PCR; primer; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ30442 standard; DNA;
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                                                                                                                                                                                                                                                                                    Sequence 24 BP; 5 A; 6 C; 5 G; 4 T; 4 other;
                                                                                                                                                                                                                                                                                                              especially Erysiphe graminis (powdery mildew).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 8; Page 35; 102pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New proteins useful for fungal infection -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Salmeron JM, Weislo LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ( SAON)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W09947552-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis thaliana.
Triticum sp.
                                                     05-AUG-2002
                                                                                 ABN99536;
                                                                                                        ABN99536 standard;
Fungus; bacteria; quantification;
                          Fungi PCR primer SEQ ID NO 108.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1999-571820/48
                                                                                                                                                                            24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOVARTIS AG. NOVARTIS-ERFINDUNGEN VERW GES MBH.
                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                            TAYGCVCTHGTCACTCAGA 6
                                                                                                                                                                                                 TTTGCGTTTGCCACTCAGA 19
                                                                                                                                                                                                                                Conservative
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                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-EP01779
                                                                                                        DNA,
                                                                                                                                                                                                                                             61.0%;
68.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     generating transgenic plants resistant to
                                                                                                         24 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
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                                                                                                                                                                                                                                           Score 12.2; DB 20;
Pred. No. 8e+03;
                                                                                                                                                                                                                                  Mismatches
 detection; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kramer. CM;
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                                                                                                                                                                                                                                Inde.s
                                                                                                                                                                                                                                                          Length
 gastroenteritis;
                                                                                                                                                                                                                                                          24;
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                                                                                                                                                                                                                                Gape
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RESULT 21
ABK29133
ID ABK29
XX ABK29
XX ABK29
XX ABK29
XX Paec:
XX Pung:
XX Pung:
XX Pung:
XX Paec
XX 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to specific fungi or bacteria detected and continued in a sample by extracting and recovering DNA from the organism continued in the sample and hybridising and amplifying the DNA sequence using continued to the species/group of species but common to all continued to the species/group of species. The method especially employs continued the species/group of species. The method especially employs continued the specifical from 225 fungal and bacterial primer and probe continued to specified bacterial. The method enables detection and continued to specification of specific fungi/fungal continued to specification of specification of specific fungi or bacterial species, useful medically for disgnosis and treatment of fungi or bacterial associated with health problems such as infections, construction associated with health problems such as infections, construction to detect and/or quantify microorganisms in the environment e.g. to establish the risk of adverse health effects (e.g. pulmonary for the contraction of specific fungional specific fungio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Paecilomyces lilacinus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ulcer; asthma; allergy; sinusitis; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK29133 standard, DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 37; Column 94; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or group of species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI, 2002-462353/49.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-APR-1998;
14-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JUN-2000; 2000US-0593012.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK29133;
                                                                                                                                                                                                                                                                                                                                                                                                 Paecilomyces lilacinus reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ussi ) us
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCGTTTGCCACTCAGAG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24 BP; 9 A; 4 C; 6 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ĠĊŢŢĠŢĠĊĄĂĊŢĊĀĠĀĠ 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vesper S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-081773P.
99US-0290990.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.2; DB 24;
Pred. No. 8e+03;
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13-JUN-2001; 2001WO-US18892

WO200196612-A2

20-DEC-2001.

Paecilomyces lilacinus

detection; bacterial detection;

probe;

primer,

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ACCESSION
VERSION
KEYWORDS
                                                                                                                                                                                                                                            REFERENCE
AUTHORS
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AI091780/c
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCUS
DEFINITION
                                                                                                                                                                          COMMENT
                                                                                                                                                                                                                                                                                                                                    SOURCE
                                  PEATURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local f
                                                                                                                                                                                         JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL; IMAGE Consortium (info@image.llnl.gov) for further 1 Trace considered overall poor quality Possible reversed clone; similarity on wrong strand Insert Length: 546 Std Error: 0.00 Seg primer: -28M13 rev2 from Amersham
                                             Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Trace considered overall poor quality
Insert Length: 439 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                 AI091780 25 bp mRNA linear EST 05-OCT-1998 ga91e02.81 Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE: 1694138 3' similar to SW:LEG7_HUMAN P47929 GALECTIN-7; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 25)
                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                     AI091780.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Wilson RK
                                                                                                                                                                                                         Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Elas of the modified pT773 vector. Library went through one round of normalization. Library constructed by M. Fatima Bonaldo."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                location/Qualifiers
organism="Homo sapiens"/
                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone_lib="Soares_pregnant_uterus_NbHPU"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'db_xref="taxon:9606"
'clone="IMAGE:471120"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           db_xref="GDB:3757122"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      50.6%;
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Pred. No. 2.1e+06;
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                                                                                                                     information.
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AUTHORS
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BM397307/c
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VERSION
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ORIGIN
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Best Local
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                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                              Seq primer: T3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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BM397307 25 bp mRNA linear EST 17-JAN-2002 5009-0-30-G06.t.1 Chilcoat/Turkewitz cDNA (large fraction)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 J. and Klobutcher, L. EST from Tetrahymena thermophila, strain CU428.1, growing cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tetrahymena thermophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BM397307.1 GI:18197360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecular Genetics and Cell Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Turkewitz AP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tetrahymena thermophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Chicago
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Conservative
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/lab_host="DH10B (ampicillin resistant)"
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/clone="IMAGE:1694138"
                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Tetrahymena thermophila"
|strain="CU428.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocation/Qualifiers
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                                    50.6%;
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Pred. No. 2.1e+06;
Score 8.6; DB 13;
Pred. No. 2.1e+06;
0; Mismatches 4;
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                                                                        Length 25;
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Query Match Best Local :

60.0%; 75.0%;

Score 12; DB 19; Pred. No. 9.6e+03;

Length 17

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RESULT 24
AAT64985
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                                               Query Match
Best Local Similarity
Matches 15; Conser
                                                                                                                               This oligonucleotide comprises forward PCR primer ROBR 27. Its sequence corresponds to bases 796-816 of a rat ob receptor (OB-R) CDNA sequence (see AAT64961). It was used with rat OB-R reverse primer ROBR 28 (see AAT64986) to amplify 1.8 kb fragments of rat genomic DNA from lean and fa/fa (fatty) Zucker rats. Subsequent digestion of the PCR product from a homozygous lean rat using restriction endonuclease MspI yielded 2 fragments of 11(0 bp and 700 bp. MspI digestion of the PCR product from a fa/fa rat yielded fragments of 950, 130 bp and 700 bp. The fa mutant OB-R gene fragments of 950, contains an A to C transversion at bp 880 that creates an extra MspI site. This allows genotype analysis of lean and
                                                                                                                                                                                                                                                                                                                                                                                                                                              25-APR-1996;
22-FEB-1996;
22-MAR-1996;
                                                                                                                                                                                                                                                                                                                             Rat wild-type and mutant ob receptor protein - useful in identification of new ligands for prevention and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat OB receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT64985;
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                                                                                                Sequence 21
                                                                                                                                                                                                                                                                                            Example 8;
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                                                                                                                                                                                                                                                                                                                                                                                              Caskey CT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ob receptor; obesity; leptin; rat; rodent; animal model; ligand; farty; fa mutation; therapy; PCR; primer; MspI; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                        (MERI ) MERCK & CO INC.
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                                                                                                                                                                                                                                                                                        Page 15; 35pp; English
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                      recertreceacteagae
                                               Conservative
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96US-0013969.
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75.0%;
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                                              Score 12; DB 18;
Pred. No. 9.9e+03;
0; Mismatches 5
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RESULT 25 AAD21460/c

RESULT 26
ABK02000
ID ABK02

ABK02000 standard; RNA; 17

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                                                                                                                                            CC like polypeptides designated as agy-96614-al and agy-969406-al and concleic acid molecules encoding such polypeptides. Polypeptides of the invention are useful for treating, preventing or ameliorating can disease, condition, or disorder which includes cancer such as a carteriosclerosis, vascular restenosis; pathology from allergens such as allergies, asthma, dermatitis, pathology from allergens such as allergies, asthma, dermatitis, dysfunction of immune system such as rheumatoid arthritis, autoimmune disease, multiple sclerosis, cancer also useful for identifying antagonists and as immunogens, for are also useful for identifying antagonists and as immunogens, for calsing antibodies which may also be used to prevent, treat or calsones a number of diseases and disorders. Polynuclocides of the invention are used to map the location of CD20/IgE-receptor like gene and related genes on chromosomes and as hybridisation probes. They are also useful in gene therapy. The present sequence is reverse transcription PCR (RT-PCR) primer used to examine the expression of human agp-96614-al sequence.
                                                                    Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel CD20/IgB-receptor like polypeptides and polynucleotides, antagonists and antibodies of the polypeptide useful for treating ameliorating or preventing diseases associated with the polypeptide e.g. cancer, asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; CD20/IgE-receptor like protein; immunoglobulin E; agp-96614-al; agp-69406-al; cancer; abnormal cell proliferation; autoimmune disease;
                                                                                                                                            Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; Page 112; 145pp; English.
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27-NOV-2000; 2000US-0723258
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infertility; gene therapy; reverse transcription PCR; RT-PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ovarian cancer; brain cancer; arteriosclerosis; vascular restenosis; rheumatoid arthritis; multiple sclerosis; allergy; dermatitis; asthma; reproductive disease; diabetes; transplant rejection; endometriosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to human CD20/immunoglobulin E (IgE)-receptor
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23
                                9 TGCCACTCAGAG 20
                                                                                       Similarity
                                                                                                                                          BP; 7 A; 7 C; 4 G; 5 T; 0 other;
                                                                    Conservative
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Pred. No.
                                                                      Mismatches
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JOURNAL COMMENT
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AUTHORS
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AU006620
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                                                                                                                                                                                                                                                                                                                                      Contact: Mitsuoki Morimyo
Genome Research Group
National Institute of Radiological Sciences
9-1, Anagawa-4-Chome, Inage-ku, Chiba, Chiba 263-8555, Japan
Email: morimyo@nirs.go.jp.
Location/Qualifiers
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22 bp mRNA linear EST 31-JU AU006620 Schizosaccharomyces pombe late log phase cDNA Schizosaccharomyces pombe cDNA clone spc00195, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schizosaccharomyces pombe
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AU006620.1
                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Morimyo, M. and Mita, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fission yeast.
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                                Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dentification of expressed sequence tags of Schizosaccharomyces
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                   schizosaccharomyces pombe was prepared by cloning cDNA into the Smal site of M13mp19 DNA and the direction of DNA sequences was not always from 5' to 3'. The cDNA data of Schizosaccharomyces pombe are available for searching on the World Wide Web. (URL, http://www.nirs.go.jp)"

1 2 6 9 11 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:1947009"
/clone_lib="NCI_CGAP_GC4"
/tissue_type="pooled_germ_cell_tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                            note="Vector: M13mp19; The cDNA library of
                                                                                                                                                                                                                              /clone_lib="Schizosaccharomyces pombe late log phase cDNA"
/sex="h minus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="Homo sapiens"
                                                                                                                                                                                                                                                                                              strain="972"
                                                                                                                                                                                                                                                                                                              'organism="Schizosaccharomyces pombe'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GI:3343079
                                                                                                                                                                                                                                                                            _xref="taxon:4896"
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                              Score 8.4; DB 9;
Pred. No. 2.6e+06;
0; Mismatches 1
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Pred. No. 2.6e+06;
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                                                           Length 22;
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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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                              TITLE
 JOURNAL
                                                                                               AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             source
                                                                                                                                                                                               GSS.
                                                                                                                                                                                                                              AZ476310
                                                                                                                                                                                                                                                                                                                                                                                                                               9;
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Mus musculus

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

1 (bases 1 to 22)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,

Dunn,D., Aoyagi,A., Barber,M., Meenen,E., Pedersen,T., Reilly

John, H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly

M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.

and Wright,D., Weises,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome Research Group
Genome Research Group
National Institute of Radiological Sciences
National Institute of Radiological Sciences
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Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AU006633 Schizosaccharomyces pombe late log phase cDNA Schizosaccharomyces pombe cDNA clone spc00210, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                               AZ476310 22 bp DNA linear GSS 04-OCT-20-
1M0294P19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0294P19 R, DNA sequence.
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 22)
Morimyo,M. and Mita,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fission yeast.
                          plasmid inserts
                                                                                                                                                                                                                                                                                                                                                             AZ476310.1 GI:10634435
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                                                                                                                                                                                                                                                                                                nouse mouse.
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l: morimyo@nirs.go.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schizosaccharomyces pombe was prepared by cloning cDNA into the Smal site of M13mp19 DNA and the direction of DN sequences was not always from 5' to 3'. The CDNA data of Schizosaccharomyces pombe are available for searching on the World Wide Web. (URL, http://www.nirs.go.jp)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="spc00210"
/clone lib="Schizosaccharomyces pombe late log phase cDNA"
/sex="h minus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'note="Vector: M13mp19; The cDNA library of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:4896"
(2000)
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Pred. No. 2.6e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence tags of Schizosaccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 22;
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RESULT 29
AAX02576/c
ID AAX025
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AC AAX025
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DE PCR pr
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Best Local S
                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                     AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent oitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX3584-AAX35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae.
PCR primer #7
                           07-MAY-1999
                                                                            AAX02576 standard;
                                                                                                                                                                                                                                                Sequence 20 BP; 6 A; 3 C; 7 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 1752; Disclosure; 1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX95488 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-357842/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-NOV-1998;
21-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
Chlamydia pneumoniae.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEST ) GENSET
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                                                                                                                                                                                                13; Conserv
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                                                                                                                                                                      GTTTGCCACTCAGAG 20
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                        (first entry)
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97FR-0014673.
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                                                                           DNA; 20
                                                                                                                                                                                                          59.0%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                               ; Score 11.8; D
; Pred. No. 1.2e
0; Mismatches
                                                                                                                                                                                                          .2e+04;
                                                                                                                                                                                                                      DB 20;
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                                                 Centromere; michrosome; vector;
                                                                                                                 20-MAR-2001
                                                                                                                                                 AAF22122;
                                                                                                                                                                              AAF22122 standard;
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 9 A; 4 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a recombinant DNA construct which comprises a functional Arabidopsis thaliana centromere. This centromeric region can be used for the production of plant artificial chromosomes (PLAC's) which can be used for the production of plants which can express
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated Arabidopsis thaliana centromeres - use production of plant artificial chromosomes for the production of plants having desirable agronomic traits
                                                                             Arabidopsis thaliana chromosome centromere associated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plant artificial chromosome; PLAC; foreign gene expression; trait; oil; herbicide; resistance; tolerance; insect; disease; stress; drought; heat; chilling; freezing; salt; moisture; oxidative; yield; food content; physical appearance; male sterility; drydown; standability; prolificacy; starch; detection; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                          method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         foreign genes. PLACs can be used to provide desirable traits such as herbicide resistance or tolerance, insect resistance or tolerance,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure, Page 77; 150pp;
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03-JUN-1997;
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                                                                                                               (first entry)
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97US-0048451.
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                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                         Score 11.8; DB 20
Pred. No. 1.2e+04;
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Arabidopsis thaliana

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JOURNAL COMMENT
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AUTHORS
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AZ511616
LOCUS
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Best Local
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                                                                                                                                                                    source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 bp DNA linear GSS 05-OCT-20
1M0356B15R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0356B15 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 20)
                                                                                                                                                                                                                                                                     Plate: 0356 row: B column: 15
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0356 row: B column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ511616.1 GI:10692932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ511616
                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                       lass: plasmid ends
igh quality sequence stop: 20.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                    /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0356B15"
                    /clone lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48.2%;
host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genome Center
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 8.2; DB 17;
Pred. No. 3.3e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GSS 05-OCT-2000
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KEYWORDS VERSION

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JOURNAL COMMENT
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AUTHORS
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AZ789409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BASE COUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DEFINITION
                                                                                                                                                                                      FEATURES
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE
                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B. W. University of Utah G. University of Utah Rm. 308, Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ789409 20 bp DNA linear GSS 16-FEB-200 2M0037H11F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0037H11 F, DNA sequence.
                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0037 row: H column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus 1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      house mouse.
                                                                                                                                                                                                                                                       Plate: 0037 row: H column: 11
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasmid inserts
                                                                                                                                                                                                          High quality sequence stop: 20.
                                                                                                                                                                                                                                                                                                                                                                                                             84112,
                                                                                                                                                                                                                                Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwb42 (gil4732114 gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli X110-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                     /organism="Mus musculus"
|strain="C57BL/6J"
|/db_xref="taxon:10090"
|/clone="Ubc2M0037H11"
|/clone=11b="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                           ocation/Qualifiers
lab_host="E. Coli strain XL10-Gold, T1-resistant,
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76.9%;
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Pred. No. 3.3e+06;
                                                                                                                                                                                                                                                                                                                Std Error: 0.00
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SOURCE

VERSION

Snoo

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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "phosphorothioate"
FEATURE:
NAME/KEY:
NAME/KEY:
NAME/KEY:
NAME/KEY:
NAME/KEY:
NAME/KEY:
NOCATION: /note= "885-905 primer"
US-08-974-549A-510

Query Match
Best Local Similarity 85.7%; Score 10.8; DB 4; Length 21;
Best Local Similarity 85.7%; Pred. No. 3.1e+03;
Matches 12; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 GTTGCTTCGGCGGG 14
Db 6 GCTTCTTCGGCGGG 19

Search completed: June 7, 2003, 09:45:18

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The present invention relates to a new polypeptide of human uridylate kinase 12, its fragment, analogue or derivative. The polypeptide of the invention and its encoding polynucleotide are applicable in diagnosis and treatment of malignant neoplasm, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases, various inflammations, sterility, pregnancy pathology, embryonic development disorders, growth and development disorders and hereditary diseases. The present nucleic acid sequence represents reverse transcriptase (RT)-PCE primer 2 that was used in the methods of the invention to isolate the coding sequence of the human kinase uridylate 12 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human uridylate kinase 12 and encoded polynucleotide, applicable diagnosis and treatment of developmental disorders, malignant tun haemopathy, HIV infection, immunological diseases and various
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Page 17; 36pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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ARESULT 34
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                 CC (I) have melting temperatures within a narrow range. The method is useful CC e.g. Salmonella, historia monocytogenes and Haemophilus influenza, fungal CC e.g. Salmonella, Listoria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus neoformans, Candida albicans and CC Aspergillus fumigautus, viruses e.g. Ticell lymphocytotrophis cirus, C Epstein-Barr virus and polio virus, and parasitic infectious agents CC selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus CC at hydroxylase deficiency, Turner Syndrome and obesity defects. CC Detecting cancer involving oncogenes, tumour suppressor genes, or genes (or genes involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC p53 gene, human papillomavirus types 16 and 18 and liver cancers. The CC method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning CC electron microscope and infrared microscope) the support at the comparise sets occurred and correlating (using a computer) identified ligation to a gressence or absence of the target nucleotide sequences. ABI82074 to CC Of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow reach minimatch, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Designing capture oligonucleotide probes complementary oligonucleotides hybridize
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-2001
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Pred. No. 1
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AI721471/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Danio rerio
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AI721471 22 bp mRNA linear EST 07-UUN-20 fc28b01.x1 Zebrafish WashU MPIMG EST Danio reric CDNA Clone fc28b01.x1 Zebrafish WashU MPIMG EST Danio reric CDNA Clone IMAGE:3722865 3' similar to TR:Q39949 Q39949 HYDROXYPROLINE-RICH PROTEIN, contains element MSR1 repetitive element ;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
Tel: 314 286 1800
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trace considered overall poor quality possible reversed clone: similarity on Seq primer: T7 ET from Amersham High quality sequence stop: 1
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Fax: 314 286 1810
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Unpublished (1998)
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                          stage embryos"
/lab_host="XL1-blue MRF"
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/clone="IMAGE:3722665"
/clone_11b="Zebrafish WashU MPIMG EST"
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                 clone UUGC2N
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AZ785027.1
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Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                  AZ785027

23 bp DNA linear GSS 16-FEB-2001
2M0028H03R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGCZM0028H03 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genomic survey sequence
                                                                                                                                                                                                                                                                10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to give a tight size distribution ( 4 kb). The \nu + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nhl@sanger.ac.uk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: nelsayed@tigr.org
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2 c
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/db_xref="taxon:5691"
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76.9%;
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                                                                                                                                                                                                                                                                Score 8.2; DB 17;
Pred. No. 3.3e+06;
0; Mismatches 3;
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Pred. No. 3.3e+06;
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RESULT 38
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AAI66253/c
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a genetically modified Saccharomyces cerevisiae in which the endogenous potassium-translocation systems (TRK1, TRK2 and TOK1) are specifically deleted and the human erg potassium ion channel (HERG) is stably integrated and expressed. This can be used to identify specific modulators of HERG, which are potentially useful as antiarrhythmic, antifibrillatory and antiinflammatory agents. The present sequence is a mutagenic primer used to disrupt the yeast TRK1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genetically modified yeast lacking endogenous potassium transport activity, useful for identifying e.g. antiarrhythmic agents, includes a functional human potassium channel -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S. cerevisiae; yeast; potassium translocation system; TRK1; TRK2; TOK1; human erg potassium ion channel; HERG; antiarrhythmic; antiinflammatory; antifibrillatory; HERG modulator; mutagenic primer; ss.
                                                                               a-cobratoxin; neurotoxin; disulphide bridge; LAS 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6 A; 7 C; 3 G; 4 T; 0 other;
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Synthetic.
Naja naja siamensis
                                                                                                                                                                     a-Cobratoxin gene amplifying primer LAS 3.
                                                                                                                                                                                                                                                                                            AAT89508;
                                                                                                                                                                                                                                                                                                                                                AAT89508 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Fig 2; 40pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cerevisiae potassium translocation gene mutagenic primer Kan-antisense
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77.8%;
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                                                                               inactivated bioactive wound healing; toxin;
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Pred. No. 1.6e+04;
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PCR primer;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          method for producing an inactivated neurotoxin with post translational modifications. A novel method for preparing a composition comprising an inactivated bioactive polypeptide comprises identifying a polypeptide having a biological activity dependent on the presence of one or more having a biological activity dependent on the presence of one or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  production of inactivated bioactive poly:peptide(s), particularly
neuro:toxin(s) - by expression of DNA encoding the polypeptide in
such a way that one or more di:sulphide bridges are not formed
                                                           alpha-cobratoxin PCR primer, LAS 3.
                                                                                                   18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a-cobratoxin. A Naja naja siamensis cDNA is used as a template used for the DNA sequencing of a-cobratoxin. This was used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 16; 32pp; English
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                                                                                                                                                                                  AAI69969 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurological diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                composition can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cardiotoxins, mellitin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         erabutoxin, a-cobratoxin and a-bungarotoxin, toxins affecting ion channels such as dendrotoxins, scorpion toxins, m-conotoxins, and sea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This primer is used for the asymmetric PCR amplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9743407-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or cell membrane-damaging toxins such as myotoxins, ellitin, and phospholipases. The inactivated neurotoxin be used for the study and treatment of viral and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Smith LA;
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77.8%;
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Pred. No. 1.6e+04;
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alpha-cobratoxin; PCR primer; anti-HIV; neurotoxin; Thailand cobra; ss.

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76 8.8 44.0 22 3	76 8.8 44.0 22 3	73 8.8 44.0 22 2	72 8.8 44.0 22 2	71 8.8. 44.0 22 2	70 8.8 44.0 22 2	69 8 8 44 0 22 2	22 0 0 44 0 0 0	1 22 0.0 14.0 0.0 15.1	0.0 44.0 66 4	0.0 44.0	64 . 88 44 0 22 1	63 8.8 44.0 22 1	62 8.8 44.0 22 1	61 8.8 44.0 22 1	1 22 O. 44 O. 0	20 0.0 14.0	EO 8 8 44 0 33 1	58 8.8 44.0 22 1	57 8.8 44.0 22 1	56 8.8 44.0 21 6		55 8 8 44 0 21 5	54 8.8 44.0 21 5	53 8.8 44.0 21 4	60 440 21 4	952 8.8 44.0 21 4	951 8.8 44.0 21 4	350 8.5 44.0 ZL 4		949 8.8 44.0 21 4	48 8.8 44.0 21 4	77 0.0 11.0 0.1		946 8.8 44.0 21 3	945 8.8 44.0 21 3	744 0.8 44.U ZI 3	044 0 0 44 0 01 0	943 8.8 44.0 21 2	942 8.8 44.0 21 2	941 8.8 44.0 21 2	210 0.0 11.0 21 2	940 88 440 21 2	939 8.8 44.U ZI Z		938 8.8 44.0 21 2	37 8.8 44.0 21 1	T T7 A. K. 0.0		935 8.8 44.0 21 1	934 8.8 44.0 21 1		933 88 44 0 21 1	932 8.8 44.0 21 1	931 8.8 44.0 21 1	0.0 44.0	930 88 440 21 1	929 8.8 44.0 21 1	928 8.8 44.0 21 1	32/ 0.0 44.0 ZT T	27 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	26 B B 44 0 20 6	8.8 44.0 20 5	24 8.8 44.0 20 4	0.0	9 44 9 99 4	8.8 44.0 20 4	8.8 44.0 . 20 4		20 8 8 44 0 30 4	8.8 44.0 20 4	18 8.8 44.0 20 4	31/ 0.0 44.0 20 4	017 0 0 110	916 8.8 44.0 20 4	0.0 HH.C	015 0 0 0 0 0 0	914 8.8 44.0 20 4	200 44.0	13 88 44 0 20 4	12 8.8 44.0 20 4	711 0.0 H	011 00 44 0 30 4	910 8.8 44.0 20 4	0.8 44.0 20.4		08 8.8 44.0 20 4	07 8.8 44.0 20 4	200 0.0 44.0 20 4	906 8 8 44 0 20 4	905 8.8 44.0 20 4	0.0 44.0 20 4	04 88 44 0 20 4
76 8.8 44.0 22	76 8.8 44.0 22 3	73 8.8 44.0 22 2	72 8.8 44.0 22 2	71 8.8. 44.0 22 2	70 8.8 44.0 22 2	69 8 8 44 0 22 2	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	0.0 11.0 CC 1	0.0 44.0 66 4	0.0 44.0	64 . 88 44 0 22 1	63 8.8 44.0 22 1	62 8.8 44.0 22 1	61 8.8 44.0 22 1	1 22 O. 44 O. 0	20 0.0 14.0	EO 8 8 44 0 33 1	58 8.8 44.0 22 1	57 8.8 44.0 22 1	56 8.8 44.0 21 6		55 8 8 44 0 21 5	54 8.8 44.0 21 5	53 8.8 44.0 21 4	60 440 21 4	952 8.8 44.0 21 4	951 8.8 44.0 21 4	350 8.5 44.0 ZL 4		949 8.8 44.0 21 4	48 8.8 44.0 21 4	77 0.0 11.0 0.1		946 8.8 44.0 21 3	945 8.8 44.0 21 3	744 0.8 44.U ZI 3	044 0 0 44 0 01 0	943 8.8 44.0 21 2	942 8.8 44.0 21 2	941 8.8 44.0 21 2	210 0.0 11.0 21 2	940 88 440 21 2	939 8.8 44.U ZI Z		938 8.8 44.0 21 2	37 8.8 44.0 21 1	T T7 A. K. 0.0		935 8.8 44.0 21 1	934 8.8 44.0 21 1		933 88 44 0 21 1	932 8.8 44.0 21 1	931 8.8 44.0 21 1	0.0 44.0	930 88 440 21 1	929 8.8 44.0 21 1	928 8.8 44.0 21 1	32/ 0.0 44.0 ZT T	27 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	26 B B 44 0 20 6	8.8 44.0 20 5	24 8.8 44.0 20 4	0.0	9 44 9 99 4	8.8 44.0 20 4	8.8 44.0 . 20 4		20 8 8 44 0 30 4	8.8 44.0 20 4	18 8.8 44.0 20 4	31/ 0.0 44.0 20 4	017 0 0 110	916 8.8 44.0 20 4	0.0 HH.C	015 0 0 0 0 0 0	914 8.8 44.0 20 4	200 44.0	13 88 44 0 20 4	12 8.8 44.0 20 4	711 0.0 H	011 00 44 0 30 4	910 8.8 44.0 20 4	0.8 44.0 20.4		08 8.8 44.0 20 4	07 8.8 44.0 20 4	200 0.0 44.0 20 4	906 8 8 44 0 20 4	905 8.8 44.0 20 4	0.0 44.0 20 4	04 88 44 0 20 4

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LENGTH: 25;
TYPE: DNA;
ORGANISM: Stachybotrys chartarum.
US-09-593-012-187
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                                                                                                                            RESULT 2
US-09-444-053-17/c
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982
983
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                                                                                                                                                                                                                                           Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
GENERAL INFORMATION:
APPLICANT: Donna T. Ward
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF NCK-2 EXPRESSION
FILE REFERENCE: RTS-0122
CURRENT APPLICATION NUMBER: US/09/444,053A
CURRENT FILING DATE: 1999-11-19
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                 Matches
                                                                                                 Sequence 17, Application US/09444053A Patent No. 6165728
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
FILE REFERENCE: HAUGLAND-1A
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                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                GTTTGCCACTCAGAG 15
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                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                            75.0%; Score 15; DB 100.0%; Pred. No. 28;
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US-09-026-039-58

US-09-417-822-27

US-08-294-312B-28

US-09-426-290-19

US-08-468-024B-28

US-09-53-231-42

US-09-538-709-193

US-08-250-849-12

US-08-250-849-12

US-08-390-850-360

US-08-588-821-35
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US-09-304-186-57
US-09-312-520-50
US-09-312-520-51
US-09-312-520-51
US-09-312-520-52
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US-08-193-627-25/c
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LENGTH: 20
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                                                                    문
                  RESULT 4
US-08-148-117A-2/c
                                                                                                                    Query Match
Best Local S
Matches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH:
                                                                                                                                                                                                                                               TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 3
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MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 90017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                 FILING DATE
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                                                                                                                                                                                              TYPE: nucleic
STRANDEDNESS:
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15; Conserve
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                                                                                                                                  Similarity
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                                                                     GAGTITGCCACTCAG 1
                                                                                             GCGTTTGCCACTCAG 18
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                                                                                                                        Conservative
                                                                                                                                                                                      linear
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93.3%;
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Pred. No. 1.2e+02;
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                                                                                                                                      Score 13.4; DB 3;
Pred. No. 1.8e+02;
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Application US/08148117A

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US-09-593-012-108
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                                                                    SOFTWARE: PatentIn version 3.0 SEQ ID NO 108 LENGTH: 24
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TYPE: DNA
ORGANISM: Paecilomyces lilacinus
-09-593-012-108
                                                                                                                                                                                                                                                                                                                                                               Sequence 108, Application US/09593012
Patent No. 6387652
                                                                                                                          PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US 60/081,773
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
                                                                                                                                                                                                                        FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
                                                                                                                                                                                                                                                                           APPLICANT: HAUGLAND, Richard
APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 421792
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,285
REFERENCE/DOCKET NUMBER: 0887-4113
REFERENCE/DOCKET NUMBER: 0887-4113
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEPHAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KENNETH H. SONNENFELD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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TITLE OF INVENTION: AG-
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PPLICANT: SCHWARZMAN, ALEXA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 22
TYPE: NUCLEOTIDE
STRANDEDNESS: SINGLE
TOPOLOGY: UNKNOWN
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STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/148,117A FILING DATE: 04-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Loca1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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82.4%;
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Pred. No. 8.
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RESULT 7
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US-08-758-306-999
                                                                                                             Matches
                                                                                                                          Query Match
Best Local
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Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REGERENCE/DOCKET NUMBER: 212/132
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/758,306
FILING DATE: December 3, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatib
OPERATING SYSTEM: IBM
SOFTWARE: PastSeq Vers
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Lyon & Ly
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APPLICANT: McSwiggen,
                                                                                                                                                                                                                                                                   TELEFAX: 10-10 TELEFAX: 67-3510
                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: lin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                         Similarity 9; Conserv
                                               UUGCCACUCAGA 12
                                                                            TIGCCACTCAGA 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3: Lyon & Lyon
633 West Fifth Street .
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                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                  (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FastSeq Version 1.5
                                                                                                                                                                                        linear
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75.0%;
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Pred. No. 8.4e+02
                                                                                                                        Score 12; DB 1;
Pred. No. 1e+03;
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                                                                                                           Mismatches
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                                                                                                           Indels
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US-08-803-346-29

Sequence 29, Application US/08803346 Patent No. 6281346

GENERAL INFORMATION:

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                                                    SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 7
                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                              APPLICANT: PREUSS, DAPHNE
APPLICANT: COPENHAVER, GREGORY
TITLE OF INVENTION: PLANT ARTIFICIAL CHROMOSOME COMPOSITIONS AND METHODS
FILE REFERENCE: ARCD:257-2
CURRENT APPLICATION NUMBER: US/09/553,231
CURRENT FILING DATE: 2000-04-19
PRIOR APPLICATION NUMBER: 60/073,741
PRIOR FILING DATE: 1998-02-05
PRIOR FILING DATE: 1998-02-05
PRIOR APPLICATION NUMBER: 60/048,451
PRIOR FILING DATE: 1997-66-03
NUMBER OF SEQ ID NOS: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                 equence 7, Application US/09553231
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0
PILING DATE: 20-FEB-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: JOANNE M. GIESSER - MERCK & CO., INC
STREET: 126 EAST LINCOLN AVENUE - P.O. BOX 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 32,838
REFERENCE/DOCKET NUMBER: 190
TELECOMMUNICATION INFORMATION:
TELEPHONE: 732-594-3046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION
NAME: GIESSER, JOANNE M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: PHILLIPS, MICHAEL SEAN
ITTLE OF INVENTION: RAT OB RECEPTORS AND NUCLEOTIDES
ITTLE OF INVENTION: ENCODING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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75.0%;
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Pred. No. 1e+03;
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic

JOTHER INFORMATION: Primer

US-09-553-231-7
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US-07-994-423-4/c
                                                                                                                         RESULT 10
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                                                                                                          US-08-421-891-4/c
                                                                       Sequence 4, Application US/08421891
Patent No. 5595969
                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: :
 APPLICANT: HAMMOND, GAPPLICANT: AVVAKUMOV, ITILE OF INVENTION: VARIABLE OF INVENTION: G
                                                                                                                                                                                                                                                                                                    STRANDEDNESS: SIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Foley & Lardner STREET: 3000 K Street, N.W., CITY: Washington, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PLICANT: AVVAKUMOV, George TLE OF INVENTION: Variants TLE OF INVENTION: Globulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/07/994,423
PILING DATE: 16-DEC-1992
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX:
                                                                                                                                                                                                                                                                                                                                                    ENGTH:
                                                                                                                                                                    20
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, Geoffrey L.
DV, George V.
Variants of Corticosteroid Binding
Globulin
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86.7%;
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Pred. No. 1
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Pred. No. 1
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Sequence 3, Apr-
No. 5989857
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/6
PTITING DATE: 10 May 1996
                                                  SOFTWARE: ASCII files
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/908,212A
FILING DATE: 07 August 1997
CLASSIFICATION: 435
                                                                                                                                                               ZIP: 55402-3397
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Mundschenk, David APPLICANT: Smith, Leonard TITLE OF INVENTION: Polypepti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FULL DATA:
CURRENT APPLICATION DATA:
US/08/421,891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 5
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                                                                                                                COMPUTER: IBM Compatible OPERATING SYSTEM: Windows-95 SOFTWARE: ASCII files
                                                                                                                                                                                                                                      STREET: 900 Second Avenue South CITY: Minneapolis STATE: Minneapola
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                       DDRESSEB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h 58.0%;
Similarity 77.8%;
14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       i: 21 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08908212A
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                                                                                                                                                                                                                                   Minnesota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              : (202)672-5300
(202)672-5399
                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                       Fredrikson & Byron, P.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-APR-1995
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                                                                                                                                                                                                                                                                                                                       Polypeptide Compositions and Methods 5
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                 08/644,399
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                                                                                                                                                                 3.50 inch, 1.44 Mb storage
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US-08-859-998-509/c
                                    ; OTHER INFORMATION: US-08-859-998-509
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Best Local :
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 599407
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                                                                                                                                                 TELEPAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
ANAME: Field, Bret E.
NAME: Field, NUMBER: 37,620
REGISTRATION NUMBER: 0909
TELECOMOUNICATION INFORMATION:
TELEPHONE: 415-322-5070
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 31,162
REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
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                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Windows95 SOFTWARE: FastSEQ for Windows Version
                                                                                              TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER:
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                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                    LENGTH: 25 base pairs
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                                                                                                                       nucleic acid
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2200 Sand Hill Road, Suite 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chenchik, Alex
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Bibilashvilli, Robert
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57.0%;
92.3%;
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                                                    oligonucleotide primer
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Pred. No. 1
 Score 11.4; DB 2; Pred. No. 2.2e+03;
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1.7e+03;
           Length 25;
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RESULT 13
US-09-225-928-509/c
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                                                                                          JS-08-164-200-9/c
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                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                   Query Match
Best Local
                                                        Sequence 9, Application US/08164200 Patent No. 5552390
                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION
                                             ENERAL INFORMATION:
                 APPLICANT:
   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                              FEATURE:
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                                                                                                                                                                                       8 TTGCCACTCAGAG 20
                                                                                                                                                                                                                                    Similarity
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OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                               OTHER INFORMATION:
NCE DESCRIPTION: SI
                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
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STREET: 2200 Sand Hill Road, Suite 100
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                Iverson,
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Bibilashvilli,
                                                                                                                                                                                                                                   57.0%;
92.3%;
Eric M.
Patrick L.
Phosphorothioate Inhibitors of Metastatic
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                                                                                                                                                                                                                                      Score 11.4;
Pred. No. 2
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HYPOTHETICAL: :
ANTI-SENSE: YE
                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
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         APPLICATION DATA:
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                                                                                                                                                                             USA
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COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.0
CURRENT APPLICATION DATA:
APPLICATION WHER: US/08/3
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/164,200
FILING DATE: December 9, 1993
CLASSIFICATION: 514
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
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NAME: Nebel, Heidi Sease
REGISTRATION NUMBER: P-3
                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE,
                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                  Philadelphia
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801 Grand Avenue Suite 3200
                                                                                                                                                                                                                                       B: Woodcock Washburn Kurtz Mackiewicz & No. 5643780ris
One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      United States
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herewith
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                                                                                                                                                                                                                                                                                                                                       Compositions and Methods for Modulating RNA Activity Through Modification of the 5' Cap Structure of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P-37,719
BER: UNMC
                      US/08/327,363
                                                                                                                      3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 11.2;
Pred. No. 2
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; TYPE: DNA
; ORGANISM: H. sapiens
US-09-328-174A-96
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                                                                                                                                                                                                                                     RESULT 17
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                                                                                                                                                                                   Patent No. 558096
                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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SEQ ID NO 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Genotyping Human Phenol Sulfotransferase TITLE OF INVENTION: (STP2)
FILE REFERENCE: 4389-6 (Formerly SEQ-16P)
CURRENT APPLICATION NUMBER: US/09/328,174A
CURRENT FILLING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 09/328,174
PRIOR FILING DATE: 1999-06-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Requence 96, Application US/09328174A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (215) 568-3439 INFORMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Guida, Marco
APPLICANT: Kurth, Janice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Kathryn Leary, Ph.D.
REGISTRATION NUMBER: 36,317
REFERENCE/DOCKST NUMBER: 16:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                             APPLICANT: HOKE, Glenn D
APPLICANT: BRADLEY, Matthews O
APPLICANT: WILLIAMS, Taffy J
APPLICANT: LEE, Che-Hung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: April 3, 19
ATTORNEY/AGENT INFORMATION:
                             CORRESPONDENCE
                                             ITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES DIRECTED ITLE OF INVENTION: AGAINST HUMAN ICAM-1 UMBER OF SEQUENCES: 15
 ADDRESSEE:
STREET: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: singl
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                                                                                                                                                                                                   Application US/08136118
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3: Naval Medical Res. & Dev. Cmd
8901 Wisconsin Ave.
                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                 ADDRESS
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81.2%;
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81.2%;
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Pred. No. 2
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Pred. No. 2.7e+03;
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                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 19;
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TELEPHONE: (202) 295-6759
TELEPAC: (202) 295-1022
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 83, Application Patent No. 6458532
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
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APPLICATION NUMBER: US 07/918,259
PILING DATE: 24-JUL-1992
ATTORNEY/AGENT INFORMATION:
NAME: Spevack, A. David
REGISTRATION NUMBER: 24,743
REFERENCE/DOCKET NUMBER: N.C. 75,77.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 20889-5606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL:
ANTI-SENSE: 1
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/091,952A
FILING DATE: 19-Apr-199
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                 Esterling, Lisa E.
TITLE OF INVENTION: Chromosomal Markers and Diagnostic
Tests for Manic-Depressive Illness
NUMBER OF SEQUENCES: 197
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                                                                                                                                                                                                           STATE: CA
                                                                                                                                                                                                                          CITY: San Francisco
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                                                                                                                                                                      ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/09091952A
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Gershon, Elliot S.
Badner, Judith A.
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                                                                                                                                                                                                                                                                                                                                                                             Berrettini, Wade
Yoshikawa, Takeo
Sanders, Alan R.
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                                                                                                                                                                                     USA
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81.2%;
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Pred. No. 2.8e+03
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PRIOR APPLICATION DATA:

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LOCATION: 1...22
OTHER INFORMATION: D18S1066 forward primer
SEQUENCE DESCRIPTION: SEQ ID NO: 83:
US-09-091-952A-83
                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 19
TYPE: DNA
ORGANISM: Human
US-09-360-416-65
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US-09-360-416-65
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US-08-393-997-1
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APPLICANT: RICHARD A. GALTI
TITLE OF INVENTION: METHODS FOR DETECTION OF ATAXIA
TITLE OF INVENTION: TELANGIECTASIA MUTATIONS
FILE REFERENCE: 510015-222
CURRENT APPLICATION NUMBER: US/09/360,416
CURRENT FILING DATE: 1999-07-23
NUMBER OF SEQ ID NOS: 143
SOPTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 65
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Patent No. 645853
                                                                                                                                                                                                                                                                                                Matches
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Best Local
                                                                                                                 Sequence 1,
                                                                                                Patent No.
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SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
GENERAL INFORMATION:
APPLICANT: Taniguchi, Tadatsugu
APPLICANT: Harada, Hisashi
TITLE OF INVENTION: Methods for Diagnosing Cancer, Precancerous
TITLE OF INVENTION: State, or Susceptibility to Other Forms of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Smith, Timothy L:
REGISTRATION NUMBER: 35,367
REFERENCE/DOCKET NUMBER: 0152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 56.0%; Score 11.2; DB 4; Length 22, Local Similarity 81.2%; Pred. No. 2.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE
                                                                                                                                                                                                                                                                                            ch 55.0%; Score 11; DB 4; 1 Similarity 100.0%; Pred. No. 3.4e+0 11; Conservative 0; Mismatches
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FILING DATE: 28-OCT-1996
APPLICATION NUMBER: PCT/US97/19381
FILING DATE: 28-OCT-1997
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                                                                                                                 Application US/08393997
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hes 0;
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS; single
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12, Application US/08927219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                atent No.
                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: 11
         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: ISM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1100 CONTROL CITY: Washington
                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
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                                                                                         CITY: Houston
STATE: Texas
COUNTRY: USA
ZIP: 77210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Steffe, Bric K.
REGISTRATION NUMBER: 36
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
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                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                      E: Arnold, Wh: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                               Bell, Graeme I.
                                                                                                                                                                                                                     Menzel, Stephen
TENTION: MUTATIONS IN THE DIABETES SUSCEPTIBILITY
TENTION: GENES HEPATOCYTE NUCLEAR FACTOR (HNF) 1 ALPHA, HNF-1BETA
JENTION: AND HNF-4ALPHA
                                                                                                                                                                                                                                                                                                                                                Yamagata,
                                                                                                                                                                                                                                                                                Horikawa, Yukio
                                                                                                                                                                                                                                                                                                  Kaisaki, Pamel
Furuta, Hiroto
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Skipping
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73.7%;
                                                                                                                                                                                                                                                                                                                  Pamela J.
                                                                                                                                                                         White & Durkee
                                                                                                                                                                                                                                                                                                                                                Kazuya
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Pred. No.
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SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS; single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                            CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Wilson, Mark B.
REGISTRATION NUMBER: 37,259
REFERENCE/DOCKET NUMBER: ARCD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512)418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENERAL INFORMATION
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APPLICATION NUMBER: US 60/028,056
PILING DATE: 02-0CT-1996
                 NFORMATION FOR SEQ ID NO: 17:
                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2
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SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60/025,719 FILING DATE: 10-SEP-1996
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                                                                                                                                                                  APPLICATION NUMBER: US/09/064,964 FILING DATE: Concurrently Herewit
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LICATION NUMBER: US 60/029,679
LICATION NUMBER: 30-0CT-1996
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VENTION: VACCINE ADJUVANTS FOR IMMUNOTHERAPY OF
                               (512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gajewski, Thomas F.
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                  ARCD: 280
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Patent No.

INFORMATION:

, Application US/08821118 5989889

APPLICANT:

Rey, Michael Golightly, Elizabeth

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; STRANDEDNESS:
; TOPOLOGY: lin
US-09-064-964-17
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US-08-821-119-21
RESULT 24
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Best Local Similarity
Matches 11; Conserv
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SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/821,119
FILING DATE: 19-MAR-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Lambiris, Blias J
REGISTRATION NUMBER: 33,728
THEORY OF THE PROPERTY OF THE PROPERT
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 41
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
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LENGTH: 23 base pairs
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 23 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                              YTGRTTYTGCCAYTCGG 23
                                                                                                                                                    TTGCGTTTGCCACTCAG 18
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Halkier, Torben
Lehmbeck, Jan
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Pred. No. 3.
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Pred. No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                        atent No. 6027906
GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
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OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/821,118
PILING DATE: 19-MAR-1997
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Diskette
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TITLE OF INVENTION:
                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                               APPLICANT: Balganesh, Tanjore S
APPLICANT: Town, Christine
TITLE OF INVENTION: No. 60279066
NUMBER OF SEQUENCES: 42
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REGISTRATION NUMBER: 33,728
REFERENCE/DOCKET NUMBER: 410
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                      COUNTRY:
                                                  APPLICATION NUMBER: US/01
FILING DATE: 10-JUL-1995
                                                                                                                                                                                                                              STATE: New York
                                                                                                                                                                                                                                                             STREET:
APPLICATION NUMBER:
                CLASSIFICATION: 435
                                                                                                                                                                                                                                                             ADDRESSEE: White & Case
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10174
                                                                                                                                                                                       TRY: United States
10036-2787
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IN 580/MAS/94
                                                                  US/08/481,435
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Pred. No. 3
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; TOPOLOGY: 11:
; MOLECULE TYPE:
US-08-481-435-18
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                                                               US-08-574-763-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/08574763 Patent No. 5736376
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Best Local Similarity
                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (212) 354-8113 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Corder, Timothy S.
REGISTRATION NUMBER: 38,414
REFERENCE/DOCKET NUMBER: UTXL
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
                                                                                                                                        TELEFAX: (512) 474-757
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: SE 9
FILING DATE: 24-NOV-1994
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/574,763 FILING DATB: Concurrently herewith CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Yanagist
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 35,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 77210-4433
               Local
                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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                                                                              TOPOLOGY:
                                                                                                                           LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13 GCGTTTGCCAC 23
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                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yanagisawa, Masashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                          : (512) 418-30
(512) 474-7577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
B: Other nucleic acid (synthethic DNA primer)
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (212) 819-8783
                                                                              linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RECOMBINANT ENDOTHELIN CONVERTING ENZYME-2 AND ITS USE IN ECE INHIBITOR SCREENING
                55.0%;
73.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SE 9404072-2
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Pred. No.
 Score 11; DB 1;
Pred. No. 3.6e+03;
0; Mismatches 5
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                                   DB 1;
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                                 Length 24;
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US-08-292-620A-10
RESULT 28
US-09-071-845-10
                                                                                                                 Query Match
Best Local Similarity
"hes 9; Conserve
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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FILING DATE: AUGUSE 17
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07
APPLICATION NUMBER: 07
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
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TITLE OF IN
TITLE OF IN
TITLE OF IN
TITLE OF IN
                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: December 7, 199
FIGNRY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                               TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
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5837542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     I: Kenneth G. Draper INVENTION: RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INVENTION: RIBOZYME TREATMENT OF INVENTION: DISEASES OR CONDITIONS INVENTION: RELATED TO LEVELS OF INVENTION: INTRACELLULAR ADHESION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTGCGTTTGCCACTCAGAG 20
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                                                                                                TTTGCCACTCAGAG 20
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633 West Fifth Street
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Dan T. Stinchcom
James McSwiggen
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                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TUMBER: US/08/292,620A
August 17, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                           January 19, 1993
UMBER: 07/989,849
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                                                                                                                                              54.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            including application described below:
                                                                                                                                                                                                                                                                                               10:
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                                                                                                                                              Score 10.8; DB 2;
Pred. No. 4.2e+03;
                                                                                                                                                             Length 15;
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                                                                                                                                Gaps
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US-08-484-956-68/c

Sequence 68, Application US/08484956

0

Patent No. 5843654

GENERAL INFORMATION:

APPLICANT:
APPLICANT:
APPLICANT:

DAHLBERG, JAMES E.
LYAMICHEV, VICTOR I.
BROW, MARY ANN D.
OLDENBURG, MARY C.

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US-09-071-845-10
                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION UNUBER: US/08/292,620
APPLICATION NUMBER: 08/008,895
FILING DATE: January 19, 1993
APPLICATION NUMBER: 07/989,849
FILING DATE: December 7, 1992
ATTORNEY/AGENT INFORMATION:
NAME: WARDURG, Richard J.
REGISTRATION NUMBER: 32,327.
REGISTRATION NUMBER: 32,327.
                                                                    Query Match
Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                              TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/071

CLASSIFT: CLASSIFT:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                      LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
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Los Angeles
California
2 UCUGCUÁCUCAGAG 15
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                                  TTTGCCACTCAGAG 20
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dan T. Stinchcomb
James McSwiggen
                                                                                                                                                                                                                                                                                   (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENTION: RIBOZYME TREATMENT OF ENTION: RIBOZYME TREATMENT OF JENETION: RIBASES OR CONDITIONS RELATED TO LEVELS OF JENTION: INTRACELLULAR ADHESION
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                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : storage
IBM Compatible
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                                                                                    54.0%;
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                                                                                                                                                                                                                                                10:
                                                                    Score 10.8; DB 3; Pred. No. 4.2e+03; 3; Mismatches 2;
                                                                                                     Length 15;
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                                                                     Gaps
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HEISLER,

LAURA

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                                                                                                                                                                                                                                  RESULT 30
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                                                                                                                                                                         Sequence 68,
Patent No. 58
                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 397-833 (NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 220 FRANCISCO CITY: SAN FRANCISCO STATE: CALIFORNIA
                                                                 APPLICANT: Kaiser, Michael W.
APPLICANT: Lyamichev, Victor I.
APPLICANT: Lyamichev, Victor I.
APPLICANT: Lyamichev, Natasha
TITLE OF INVENTION: Cleavage Of Nucleic Acid Using
TITLE OF INVENTION: Thermostable FEN-1 Endonucleases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: FO TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 0 FILING DATE: 07-DEC-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PILING DATE: 04-JUN-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOPTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: ...
STRANDEDNESS: b...
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                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: UNITED STATES OF AMERICA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/254,359
                  DDRESSEE
                                                                                                                                                                                                                                                                                      19 TTTGCCACACTGAG 6
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12, Conserv
                                                                                                                                                                         8, Application US/08757653 .
5843669
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                                                                                                                                                                                                                                                                                                                        TTTGCCACTCAGAG 20
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220 MONTGOMERY STREET, SUITE 2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (415) 397-8338
220 Montgomery Street, Suite 2200
                                                                                                                                                                                                                                                                                                                                                           Conservative
                Medlen & Carroll, LLP
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85.7%;
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Pred. No. 4.4e+03;
0; Mismatches 2;
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Patent No. 58517
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MOLECULE TYPE: DNA (genomic):-08-757-653-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,952
FILING DATE: 07-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: INGOLIA, Diane E.
REGISTRATION NUMBER: 40,027
REFERENCE/DOCKET NUMBER: FORE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 397-8338 INFORMATION FOR SEQ ID NO:
                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/078,471
FILING DATE: 15-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Los Angeles
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
NAME: Reiter, Stephen E. REGISTRATION NUMBER: 31,192 REFERENCE/DOCKET NUMBER: P4: LECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                           E: Pretty, Schroeder, Brueggemann & Clark 444 South Flower Street, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08117952
                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Evans, Glen A.
Smith, Michael W.
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85.7%;
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                  P41 9423
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Pred. No. 4.
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                                                                                                                                                                                                                 Version #1.25
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Sequence 29, Application US/09428696
Patent No. 6165789
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF HURNP A1
FILE REPERENCE: RTS-0111
CURRENT APPLICATION NUMBER: US/09/428,696
CURRENT FILING DATE: 1999-10-27
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                   RESULT 33
US-09-428-696-29/c
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-358-685-23
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US-08-117-952-652
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SEQ ID NO 23
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Best Local Similarity
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 Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                             SEQ ID NO 29
LENGTH: 20
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                                                         OTHER INFORMATION: Antisense Oligonucleotide -09-428-696-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/358,685
CURRENT FILING DATE: 1999-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
TITLE OP INVENTION: ANTISENSE MODULATION OF SHP-1 EXPRESSION
FILE REFERENCE: RTS-0081
                                                                                    TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
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TOPOLOGY: lir
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  Conservative
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              54.0%;
85.7%;
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Score 10.8; DB 4; Length 20; Pred. No. 4.4e+03; 0; Mismatches 2; Indels
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Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 10.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                 8; DB 3;
4.4e+03;
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US-08-520-946-68/c
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LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 68, Application US/08520946 Patent No. 6372424
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APPLICANT: RUSCHOff, JOSEf
TITLE OF INVENTION: IMPROVED METHOD OF PRIMER-EXTENSION PREAMPLIFICATION
TITLE OF INVENTION: PCR
FILE REFERENCE: 4802
CURRENT APPLICATION NUMBER: US/09/270,933
CURRENT FILING DATE: 1999-03-16
EARLIER APPLICATION NUMBER: DE 198 13 317.0
EARLIER APPLICATION NUMBER: DE 198 13 317.0
EARLIER FILING DATE: 1998-03-26
NUMBER OF SEQ ID NOS: 17
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PATENTIN RC-BOSMS-DOS
SOPTWARE: PATENTIN RATA:
APPLICATION NUMBER: US/08/520,946
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FORS-01756
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                             ZIP: 94104
COMPUTER READABLE FORM;
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: RAPID DET TITLE OF INVENTION: PATHOGENS UMBER OF SEQUENCES: 160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BROW, MARY ANN D.
APPLICANT: LYAMICHEV, VICTOR I.
APPLICANT: OLIVE, DAVID M.
                                                                                                                                                                                                                                                                                                                                                         STATE:
                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                               STREET:
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SAN FRANCISCO
: CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                   UNITED STATES OF AMERICA
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85.7%;
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                                                                                                                                                                    US/08/520,946
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                                              FORS-01756
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                                                                                                                                                                                                            Version #1.25
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Query Match
Best Local Similarity
Marches 12; Conserva
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                                                                       US-08-415-818-8
Query Match
Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                     TELBFAX: 908-594-4720 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/415,818
FILING DATE:
CLASSIFICATION: 435
                                                                                                     STRANDEDNESS: single TOPOLOGY: linear HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/383,746
PILING DATE: 03-FBB-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Linemeyer, David L.
APPLICANT: MacNeil, Douglas J.
                                                                                                                                                                                                                                      REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19390
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3462
                                                                                     ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: P.O. CITY: Rahway STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                    TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                      NAME: Appollina, Mary A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 07065
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P.O. Box 2000, 126 E. Lincoln Ave.
                                                                                                                                                                                                                        908-594-4720
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Tan, Carina P.
Weinberg, David H.
VENTION: NEUROPEPTIDE Y RECEPTOR
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      Conservative
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85.7%;
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     0; Mismatches
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Pred. No. 4.
                    Score 10.8; DB 1; Length 21; Pred. No. 4.5e+03;
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     Gaps
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            RESULT 38
PCT-US96-01444-8/c
Sequence 8, Application PC/TUS9601444
; GENERAL INFORMATION:
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                                                                                                                                          Matches
                                                                                                                                                                                                              Query Match
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INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/415,818
FILING DATE: 03-APR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/3
FILING DATE: 03-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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STREET: F.
STREET: NAMES
NAMES OF THE STREET: S
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                                                                                                                                                                        Local
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                      l Similarity
12; Conservat
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                                            TIGCGTTIGCCACT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Mary A. Appollina
P.O. Box 2000, 126 E. Lincoln Ave.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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PENTION: NEUROPEPTIDE Y RECEPTOR
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                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     linear
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85.7%;
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                                                                                                                                                                                Score 10.8; DB 2;
Pred. No. 4.5e+03;
                                                                                                                                                                                                              Length 21;
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Cascieri, Margaret A. Linemeyer, David L.

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PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/383,746

PILING DATE: 03-FEB-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/415,818

PILING DATE: 03-APR-1995

ATTORNEY/AGENT INFORMATION:

NAME: APPOILIAID, MALY A.

REGISTRATION UNMBER: 19,087

REFERENCE/DOCKET NUMBER: 19390Y

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                    ENERAL INFORMATION:
APPLICANT: Cheng, Suzanne
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LENGTH: 21 base pairs
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                                                                                                                                   NUMBER OF SEQUENCES: 3
                                                                                                                                                       TITLE OF INVENTION: Methods and Reagents for the Polymerase TITLE OF INVENTION: Chain Reaction Amplification of Long DNA Sequences NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE:
OMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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                                                                                                     340 Kingsland Street
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VENTION: NEUROPEPTIDE Y RECEPTOR
                                                  U.S.A.
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Strader, Catherine D
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                                                                                                                     Hoffmann-La Roche Inc
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O. Box 2000, 126 E. Lincoln Ave.
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Pred. No. 4.5e+03;
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Search completed: June 7, 2003, 09:45:29 Job time: 53.1818 secs

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                                                                                                                              US-09-358-685-2
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                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 47
SEQ ID NO 2
LENGTH: 22
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09358685
Patent No. 6121047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                            APPLICANT: C. Frank Bennett
APPLICANT: Lex M. COWBERT
TITLE OF INVENTION: ANTISENSE MODULATION OF SHP-1 EXPRESSION
FILE REFERENCE: RTS-0081
CURRENT APPLICATION NUMBER: US/09/358,685
CURRENT FILING DATE: 1999-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (510) 814-29
TELEFAX: (510) 814-2977
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                              OTHER INFORMATION: PCR Primer
                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
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NAME: Petry Ph.D., Douglas A.
REGISTRATION NUMBER: 35,321
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TELECOMMUNICATION INFORMATION:
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                                                                12;
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                                                                               Similarity
                              GTTTGCCACTCAGA 19
GTTTGCGACTCTGA 20
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85.7%;
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Pred. No. 4.5e+03
                                                              Score 10.8; DB 3;
Pred. No. 4.5e+03;
0; Mismatches 2
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                                                                                                DB 3;
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                                                                                             Length 22;
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Title:
Perfect score:
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; OTHER INFORMATION:
US-10-080-959A-2
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RESULT 2
US-10-080-959A-4
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APPLICANT: Buttner, Mark P.
APPLICANT: Buttner, Mark P.
TITLE OF INVENTION: Method for Detection of Stachybotrys chartarum in Pure Culture of ILLE REFERENCE: 0001-00001
FILE REFERENCE: 0001-00001
CURRENT APPLICATION NUMBER: US/10/080,959A
CURRENT FILING DATE: 2002-05-16
PRIOR APPLICATION UNMBER: US 60/280,712
PRIOR FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 5
SEQ ID NO
TYPE: DNA
ORGANISM: Stachybotrys chartarum
PEATTER:
                                                                                        Query Match
Best Local S
Matches 20
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NAME/KEY:
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20; Conserv
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                                              TTTGCGTTTGCCACTCAGAG 20
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US-10-096-718-30

US-08-591-466B-122

US-10-056-414-336

US-10-056-414-337

US-10-056-414-337

US-10-056-414-338
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US-09-864-785-303

US-09-864-785-560

US-09-864-785-1555

US-09-864-785-2121

US-10-079-625-27

US-10-079-625-27
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US-09-827-998-1580

US-09-827-998-1581

US-09-989-731-237

US-09-989-731-237

US-09-989-732-237

US-09-991-073-237

US-09-991-073-237
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US-09-993-604-237
US-09-990-456-237
US-09-989-721-237
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US-09-274-553D-273
US-09-829-855-172
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Pred. No. 0.2
0; Mismatches
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                                                                                                                9; Length 20
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Sequence 1576, Ap
Sequence 1579, Ap
Sequence 1579, Ap
Sequence 1581, Ap
Sequence 237, App
Sequence 30, Appl
Sequence 30, Appl
Sequence 316, App
Sequence 317, App
Sequence 273, App
Sequence 273, App
Sequence 318, App
Sequence 273, App
Sequence 373, App
Sequence 374, App
Sequence 3755, App1
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APPLICANT: McSwiggen, Jim
FILE Reference: McSwiggent for Inhibiting Hepatitis B Virus Replication
CURRENT FILING DATE: 2001-12-31
CURRENT FILING DATE: 2001-12-31
PRIOR APPLICATION NUMBER: US 09/882,712
PRIOR APPLICATION NUMBER: US 09/531,025
PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR FILING DATE: 2000-08-09
PRIOR APPLICATION NUMBER: US 09/636,385
PRIOR FILING DATE: 1994-00-00-09
PRIOR APPLICATION NUMBER: US 09/696,347
PRIOR APPLICATION NUMBER: US 08/193,627
PRIOR APPLICATION NUMBER: US 08/193,627
PRIOR APPLICATION NUMBER: US 08/433,993
PRIOR APPLICATION NUMBER: US 08/433,993
PRIOR APPLICATION NUMBER: US 08/434,504
PRIOR APPLICATION NUMBER: US 08/434,504
PRIOR APPLICATION NUMBER: US 08/434,504
PRIOR FILING DATE: 1995-05-04
PRIOR FILING DATE: 1995-05-04
PRIOR FILING DATE: 1995-10-08
NUMBER OF SEQ ID NOS: 6586
SOFTWARE: PatentIn version 3.0
SEQ ID NO 25
LEGGTH: 17
TYPE: RNA
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; OTHER INFORMATION:
US-10-080-959A-4
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                                                                                                           ; TYPE: RNA
; ORGANISM: Hepatitis
US-09-877-478-25
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Best Local Similarity 100.
Matches 17; Conservative
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SEQ ID NO 4
LENGTH: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cruz-Perez, Patricia
APPLICANT: Butther, Mark P.
APPLICANT: Butther, Mark P.
TITLE OF INVENTION: Method for Detection of Stachybotrys chartarum in Pure Culture
TITLE OF INVENTION: Samples Using Quantitative Polymerase Chain Reaction
FILE REFERENCE: 0001-00001
CURRENT APPLICATION NUMBER: US/10/080,959A
CURRENT FILING DATE: 2002-05-16
PRIOR APPLICATION NUMBER: US 60/280,712
PRIOR FILING DATE: 2001-03-29
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Query Match
Best Local Similarity 93.3
Warches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
TYPE: DNA
ORGANISM: Stachybotrys chartarum
FEATURE:
NAME/KEY:
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Blatt, Larry
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                                                   Score 13.4; DB 9;
Pred. No. 9.4e+02;
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SEQ ID NO 10
LENGTH: 23
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APPLICANT: PFIZER INC.
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US-09-780-533A-2611
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                                                         RESULT 6
US-09-819-097-24/c
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Best Local Similarity
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ORGANISM: Human
-09-972-467-10
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LENGTH: 17
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Best Local Similarity
                                                                                                                                                                                     Matches
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CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
                           Sequence 24, Application US/09819097 Patent No. US20020106347A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 6679
SOFTWARE: PatentIn version 3.0
APPLICANT:
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Douglas, Andrea M.
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Pred. No. 1
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Pred. No. 4
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TITLE OF INVENTION: CYTOKINES AND THEIR USE IN TREATMENT AND/OR PROPHYLAXIS
TITLE OF INVENTION: OF BREAST CANCER
FILE REFERENCE: 11375Z
CURRENT APPLICATION NUMBER: US/09/819,097
CURRENT FILING DATE: 2001-03-05
CURRENT FILING DATE: 09051,939
PRIOR APPLICATION NUMBER: 09/051,939
PRIOR FILING DATE: 1998-10-16
NUMBER OF SEQ ID NOS: 28
NUMBER OF SEQ ID NOS: 28
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                                                                                                                                                                 RESULT 8
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LENGTH: 23
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Welcher,
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                                                                                                              Sequence 2000, Application US/09780533A Publication No. US20030060611A1
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Best Local Similarity
                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/821,821
CURRENT FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: US 09/723,258
PRIOR FILING DATE: 2000-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Welcher, Andrew A.
APPLICANT: Calzone, Frank J.
TITLE OF INVENTION: CD20/IgE-Receptor Like Molecules and Uses Thereof FILE REFERENCE: 01017/36938A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/193,728 PRIOR FILING DATE: 2000-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 25
                                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc
PPLICANT: Habberil, rece
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                  23 recenerciación 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGCGTTTGCCACTCAGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09821821
                                                                                                                                                                                                                                                       TGCCACTCAGAG 20
                                                                  Blatt, Larry
                     McSwiggen, Jim
Chowrira, Bharat
Haeberli, Pete
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                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61.0%;
                                                                                                                                                                                                                                                                                                        60.0%; Score 12;
100.0%; Pred. No.
                                                      Jim
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Pred. No. 4.2e+03;
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                                                                                                                                                                                                                                                                                                               DB 10;
5.3e+03;
                                                                                                                                                                                                                                                                                                                               Length 23;
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; PILE REFERENCE: MBHBOO,878-A (400/011)
; CURRENT APPLICATION NUMBER: US/09/780,533A
; CURRENT FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: US 60/181,797
; PRIOR PILING DATE: 2000-02-11
; NUMBER OF SEQ ID NOS: 6679
; SEQ ID NO 2000
; SEQ ID NO 2000
; LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                 RESULT 10
US-09-368-834-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) ORGANISM: Homo sapiens
US-09-780-533A-2000
                                                                                                                                                                                                                                                                                              Sequence 3, Application US/09368834
Patent No. US20020068317A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOPTWARE: PatentIn version 3.1
SEQ ID NO 14
LENGTH: 22
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14, Application US/09996956 Patent No. US20020155463A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Prostate Polynucleotides and Uses
FILE REFERENCE: 9U 301 R1
CURRENT APPLICATION NUMBER: US/09/956,956
CURRENT FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: 60/250,354
PRIOR FILING DATE: 2001-12-01
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: OriGene Technologies, Inc
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Local Similarity 60.0%;
les 9; Conservation
              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 i
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows-95
SOFTWARE: ASCII files
                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                     TITLE OF INVENTION: Polypeptide Compositions and Methods NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 TGCGTTTGCCACTCA 17
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Similarity 86.7%;
                                                                                                                                                      ADDRESSEE: Fredrikson & Byron, P.A. STREET: 900 Second Avenue South CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UGCUUUGGCCACUCA 16
                                                                                                                                  STATE: Minnesota
                                                                                                     ZIP: 55402-3397
                                                                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
   APPLICATION DATA:
                                                                                                                                                                                                                                                                            Mundschenk, David
                                                                                                                                                                                                                                                               Smith,
                                                                                                                                                                                                                                                           Leonard
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Pred. No. 6
                                                               3.50 inch, 1.44 Mb storage
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                                                             RESULT 12
US-08-844-215-30
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          Sequence 30, Application US/08844215 Patent No. US20020016445A1 GENERAL INFORMATION:
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/940,185
CURRENT FILING DATE: 2001-08-27
PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
SOFTWARE: Patentin version 3.1
SEQ ID NO 2248
LENGTH: 24
                                                                                                                                              US-09-940-185-2248
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                                                         Query Match
Best Local Similarity 77.8
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2248, Application US/09940185 Publication No. US20030096239A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A-69605-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Gunderson, lappLICANT: Chee, Mark
                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                              OTHER INFORMATION: Computer Generated Probe Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 15050.1,2.2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 347-7088
TELEPAX: (612) 347-7077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/908,212
18
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                          TTTGCGTTTGCCACTCAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTTGCGTTTGCCACTCAG 18
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TITGCGTTTTCCAACCGG
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APPLICATION NUMBER: 08/644,399
FILING DATE: 10 May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Goldman, Philip M. REGISTRATION NUMBER: 31,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/368,834 FILING DATE: 05-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
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                                                                                     58.0%;
77.8%;
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                                                                                     Score 11.6; DB 9;
Pred. No. 8.7e+03;
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Pred. No. 8.6e+03
                                                                      Mismatches
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                                                                                                   Length 24;
                                                                    Indels
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PERSSON, MATS AXEL

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APPLICANT: Steinberger, Peter
TITLE OF INVENTION: METHODS AND COMPOSITIONS USEFUL FOR
TITLE OF INVENTION: INHIBITING CCR5-DEPENDENT INFECTION OF CELLS BY HIV-1
FILE REFERENCE: TSR1 728.1
CURRENT APPLICATION NUMBER: US/09/913,238
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: PCT/EPO 0/12419
PRIOR APPLICATION NUMBER: US/09/9653
PRIOR APPLICATION NUMBER: US 60/169,653
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 85
SOPTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 53
LENGTH: 24
TYPE: DNA
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 13
US-09-913-238-53
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US-08-844-215-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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OTHER INFORMATION: Synthesized 09-913-238-53
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APPLICANT: Steinberger, Pet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (650) 325-782
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                      ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NIMBER
                                       PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 38,548
REFERENCE/DOCKET NUMBER: 80
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 325-7812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: ROBINS & ASSOCIATES
STREET: 90 MIDDLEFIELD ROAD, SUITE 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 19-APR-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ALLANDER FITLE OF INVENTION: FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TUMBER OF SEQUENCES:
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TOPOLOGY: linear
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FILING DATE: 17-APR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/635,109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: MCCRACKEN, THOMAS P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTAGCGTTTGCCA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTTGCGTTTGCCA 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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HUMAN MONOCLONAL ANTIBODIES SPECIFIC FOR
HEPATITIS C VIRUS (HCV) E2 ANTIGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57.0%;
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APPLICANT: Higher, Jim
APPLICANT: Chowrira, Bharat
APPLICANT: Hasberli, Pete
TITLE OF INVENTION: Method and Reagent for the Inhibition of NOGO Gene
FILE REFERENCE: MBHB00,878-A (400/011)
CURRENT APPLICATION UNMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
FRIOR APPLICATION NUMBER: US 60/181,797
FRIOR APPLICATION NUMBER: US 60/181,797
FRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
SOFTWARE: Patentin version 3.0
SEQ ID NO 505
LENGTH: 17
밁
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                                                                                                              ; TYPE: RNA; Homo sapiens US-09-780-533A-505
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Best Local S
Matches 12
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10761
LENGTH: 25
TYPE: DNA
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                                                         Matches
                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                      Local Similarity hes 8; Conserv
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ucudcuuugdccacuc 17
                           TTTGCGTTTGCCACTC
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                                                        Conservative
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                                                                  56.0%; Score 11.2;
50.0%; Pred. No. 1.
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92.3%;
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Pred. No. 1.1e+04;
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Pred. No. 1.
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                                                        Mismatches
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                                                                  .4e+04;
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                                                                                Length 17;
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US-09-930-423-1153/c
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US-10-206-839-96
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US-09-930-423-1153
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APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for TITLE REFERENCE: MBHB00, 918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930, 423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
          Sequence 158, Application US/10060301

Publication No. US20020182622A1

GENERAL INFORMATION:
APPLICANT: NAKAMURA, YUGUKe et al.
TITLE OF INVENTION: A METHOD FOR SNP (SINGLE NUCLEOTIDE POLYMORPHISM) TYPING
FILE REFERENCE: 1254-0195p
CURRENT APPLICATION NUMBER: US/10/060,301
CURRENT FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: JP 2001-25700
PRIOR FILING DATE: 2001-02-01
PRIOR FILING DATE: 2001-02-01
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Best Local Similarity
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APPLICANT: Kurth, Janice
TITLE OF INVENTION: Genotyping Human Phenol Sulfotransferase
TITLE OF INVENTION: (STP2)
FILE REFERENCE: 4389-6 (formerly SEQ-16P)
CURRENT APPLICATION NUMBER: US/10/206,839
CURRENT FILING DATE: 2002-07-26
PRIOR APPLICATION NUMBER: 09/328,174
PRIOR FILING DATE: 1999-06-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 19
TYPE: DNA
ORGANISM: H. sapiens
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NUMBER OF SEQ ID NOS: 200
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ENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                      Local
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81.2%;
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Pred. No. 1.4e+04;
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Pred. No. 1.
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CURRENT APPLICATION NUMBER: US/09/808,680
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/194,230
PRIOR FILING DATE: 1999-02-24
PRIOR FILING DATE: 1999-02-24
PRIOR FILING DATE: 1997-04-29
PRIOR FILING DATE: 1997-04-29
PRIOR APPLICATION NUMBER: 08/653,653
PRIOR APPLICATION NUMBER: 08/440,740
PRIOR PILING DATE: 1995-05-12
                                                                                                                                                                                                                                                                                                                                                                                              US-09-808-680-10/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: NAKAMURA, YUSUKe et al.
TITLE OF INVENTION: A METHOD FOR SNP (SINGLE NUCLEOTIDE POLYMORPHISM) TYPING
FILE REFERENCE: 1254-0195p
CURRENT APPLICATION NUMBER: US/10/060,301
CURRENT FILING DATE: 2002-02-01
CURRENT FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: JP 2001-25700
PRIOR FILING DATE: 2001-02-01
NUMBER OF SEQ ID NOS: 200
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 160
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Best Local (
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OTHER INFORMATION:
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TYPE: DNA
ORGANIEM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                       APPLICANT: Condon, Thomas P
TITLE OF INVENTION: Compositions and Methods for Antisense Inhibition
TITLE OF INVENTION: Protein Translation
                                                                                                                                                                                                                                                                                             APPLICANT: Baker, APPLICANT: Bennet
                                                                                                                                                                                               FILE REFERENCE: ISPH-0557
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ORGANISM: Artificial Sequence
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nes 13; Conserv
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Reverse Primer for SNP ID 79
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81.2%;
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pred. No. 1.4e+04;
0; Mismatches 3;
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Pred. No. 1.4e+04;
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RESULT 22
US-10-175-225-65
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; NAME/KEY: misc_feature
; LOCATION: (1)...(6)
; OTHER INFORMATION: The monomer thegog may be repeated from US-10-117-108A-27
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              Sequence 65, Application US/10175225
Publication No. US20030082582A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 27
LENGTH: 12
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic oligonucleotide
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Best Local &
                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/117,108A CURRENT FILING DATE: 2002-04-08 PRIOR APPLICATION NUMBER: US 60/282,491 PRIOR FILING DATE: 2001-04-10 NUMBER OF SEQ ID NOS: 80 SOFTMARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BARNETT, Graeme R.
TITLE OF INVENTION: LINEAR NUCLEIC ACID AND SEQUENCE THEREFOR FILE REFERENCE: 37955-0004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 07/568,366
PRIOR FILING DATE: 1990-08-16
NUMBER OF SEQ ID NOS: 15
APPLICANT: Richard A. Gatti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KACHAB,
APPLICANT: BARNET
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TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
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APPLICATION NUMBER: 07/939,855
FILING DATE: 1992-09-02
APPLICATION NUMBER: 07/567,286
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APPLICATION NUMBER: 07/927,506
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                                                                                                                                                                                                55.0%; Score 11; DB 9;
100.0%; Pred. No. 1.7e+0:
tive 0; Mismatches
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Pred. No. 1.
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Blatt, Larry
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Chowrita, Bharat
APPLICANT: Haeberli, Pete
TITLE OF INVENTION: Mcthod and Reagent for the Inhibition of
FILE REFERENCE: MBHB00, 878-A (400/011)
CURRENT APPLICATION NUMBER: US/09/780,533A
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: US 60/181,797
PRIOR FILING DATE: 2000-02-11
NUMBER OF SEQ ID NOS: 6679
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; OTHER INFORMATION: Sense PCR primer US-09-801-968-5
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US-10-175-225-65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: PP-17150.001/201130.40901
CURRENT APPLICATION NUMBER: US/09/801,968
CURRENT FILING DATE: 2001-03-07
NUMBER OF SEQ ID NOS: 46
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 20
                                                                                                                                                                                                                                             GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Kavanaugh, W. Michael
TITLE OF INVENTION: HUMAN FGF-23 GENE AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/175,225
CURRENT FILING DATE: 2002-08-23
PRIOR APPLICATION NUMBER: US 09/360,416
PRIOR FILING DATE: 1999-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHODS FOR DETECTION OF ATAXIA TITLE OF INVENTION: TELANGIECTASIA MUTATIONS FILE REFERENCE: UC078.001DV1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 143
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US-10-087-082-2/c
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                                                                                                         ) OTHER INFORMATION: Antisense Oligonucleotide US-09-865-866-22
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Patent No. US/20020160404A1
GENERAL INFORMATION:
APPLICANT: Dietmaler, Wolfgang
APPLICANT: Ruschoff, Josef
                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/865,866
CURRENT FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 173
SEQ ID NO 22
LENGTH: 20
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Matches
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 20
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Best Local &
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LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                               Matches
                                                           Query Match
Best Local Similarity
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APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOLIPASE A2, GROUP IIA (SYNOVIAL)
FILE REFERENCE: RTS-0221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/087,082
CURRENT FILING DATE: 2002-02-28
PRIOR APPLICATION NUMBER: DE 198 13 317.0
PRIOR FILING DATE: 1998-03-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: IMPROVED METHOD OF PRIMER-EXTENSION PREAMPLIFICATION PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                     ORGANISM: Artificial Sequence PEATURE:
                                                                                                                                                                   TYPE: DNA
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                                             12;
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Pred. No. 2.3e
3; Mismatches
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Pred. No. 2.3e+04;
                                             Score 10.8; DB 9;
Pred. No. 2.3e+04;
0; Mismatches 2;
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2.3e+04;
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US-09-865-866-23
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Best Local
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RESULT 28
US-09-940-925A-68/c
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CURRENT APPLICATION NUMBER: US/09/865,866
CURRENT FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 173
SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 68, Application No. US2003
Publication No. US2003
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                   TELEFAX: (415) 397-8
INFORMATION FOR SEQ ID NO: 68:
                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FOI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PETENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BROW, MARY ANN D.
LYAMICHEV, VICTOR I.
OLIVE, DAVID M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: RAPID DETECTION AND IDENTIFICATION OF
                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
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1 Similarity 85.7%;
12; Conservation
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                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/940,925A FILING DATE: 10-Jun-2002 CLASSIFICATION: <Unknown>
                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UNITED STATES OF AMERICA
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                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
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o. US20030054338A1
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(genomic)
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Pred. No. 2.
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Best Local S
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CURRENT FILING DATE: 2001-10-23
NUMBER OF SEQ ID NOS: 179
SEQ ID NO 59
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6, Application US/10011119A Patent No. US20020150928A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 59, Application US/10001076 Publication No. US20030096775A1 GENERAL INFORMATION:
                                                                       Query Match
Best Local (
                                                                                                                                                                                                                        SEQ ID NO 6
                                                                                                                                                                                                                                                                                                  PILE REPERENCE: P/2432-45
CURRENT APPLICATION NUMBER: US/10/011,119A
CURRENT FILING DATE: 2001-11-13
                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 09/605,611
PRIOR FILING DATE: 2000-06-28
                                                                                                                    10-011-119A-6
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF
FILE REFERENCE: RTS-0329
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APPLICANT: Lundin, Toma
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TYPE: DNA
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                                                                                                                            PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: Construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                             TYPE: DNA
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ITLE OF INVENTION: DNA-EMBEDDING MEDIUM AND METHOD OF USE
                                                                     Local
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Pred. No. 2
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; Sequence 15, Application US/10135629
; Publication No. US20030073657A1
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Best Local Similarity
Matches 12; Conserv
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LENGTH: 24
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Publication No.
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SEQ ID NO 15
LENGTH: 21
TYPE: DNA
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PRIOR FILING DATE: 2000-06-15
NUMBER OF COLUMBER 2000-06-15
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TITLE OF INVENTION:
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: DE 10121225.0
PRIOR FILING DATE: 2001-04-30
PRIOR APPLICATION NUMBER: US 60/323,348
PRIOR FILING DATE: 2001-09-18
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CURRENT FILING DATE: 2001-06-15
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ORCANISM: Artificial Sequence
FEATURE:
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PHICANT: Reinhard, Christoph
PHICANT: Jefferson, Anne Bennett
THE OF INVENTION: POLYNUCLEOTIDES RELATED TO COLON CANCER
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Pred. No. 2.3e
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Pred. No. 2.
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2.3e+04;
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; ORGANISM; Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Synthetic Oligonuclectide
US-10-215-112-665
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                                                                                                                                             RESULT 35
                                                                                                                              US-10-215-112-79
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 665
LENGTH: 25
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Publication No.
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                                                   Sequence 791, Application US, Publication No. US20030082590 GENERAL INFORMATION:
APPLICANT: Michael Mittmann
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Best Local
                                                                                                                                                                                                                                                                           Matches
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SOPTWARE: PatentIn version 3.1
SEQ ID NO 2382
LENGTH: 24
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ORGANISM: Artificial Sequence
FEATURE:
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ITLE OF INVENTION: Method of Genetic Analysis of Probes: ITLE OF INVENTION: Test3
ILE REFERENCE: 3119
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o. US20030082596A1
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85.7%;
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85.7%;
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Pred. No. 2.3e
0; Mismatches
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Synthetic Oligonucleotide US-10-215-112-2803
                                ; OTHER INFORMATION: Synthetic Oligonucleotide US-10-215-112-2929
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SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 791
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                                                                                                                 CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
RUMBER OF SEQ ID NOS: 14936
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 2929
LENGTH: 25
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APPLICANT: Michael Mittmann
TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 2803
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Best Local 9
Query Match
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CURRENT FILING DATE: 2002-08-08
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Michael Mittmann
TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 25
TYPE: DNA
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                   ORGANISM: Artificial Sequence
                                                                                                     TYPE: DNA
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US20030082596A1
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85.7%;
   54.0%; Score 10.8; DB 9;
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Pred. No. 2.3e+04;
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PILE REFERENCE: 3119
CURRENT APPLICATION NUMBER: US/10/215,112
CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOPTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 8357
LENGTH: 25
TYPE: DNA
ORGANISM: Artificial Sequence
RESULT 40
US-10-215-112-12458
US-10-215-112-12458
/ Sequence 12458, Application US/10215112
/ Publication No. US20030082596A1
/ GENERAL INFORMATION:
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Best Local Similarity
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US-10-215-112-11650/c
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US-10-215-112-8357
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Publication No. US20030082596A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-08-08
NUMBER OF SEQ ID NOS: 14936
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 11650
LENGTH: 25
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Best Local Similarity 85.7%;
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Matches 12; Conservative
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Publication No. US20030082596A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Michael Mittmann
TITLE OF INVENTION: Method of Genetic Analysis of Probes:
TITLE OF INVENTION: Test3
FILE REFERENCE: 3119
                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENERAL INFORMATION:
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ITLE OF INVENTION: Method of Genetic Analysis of Probes:
ITLE OF INVENTION: Test3
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                                                                                                                       24 CITITGCTACTCAG 11
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Pred. No. 2.3e+04;
0; Mismatches 2;
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; APPLICANT: Michael Mittmann; ITITLE OF INVENTION: Method of Genetic Analysis of Probes: ITITLE OF INVENTION: Test3
; FILE REFERENCE: 3119
; CURRENT APPLICATION NUMBER: US/10/215,112
; CURRENT FILLING DATE: 2002-08-08
; NUMBER OF SEQ ID NOS: 14936
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 12458
; LENGTH: 25
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FCATURE:
; OTHER INFORMATION: Synthetic Oligonucleotide
US-10-215-112-12458

QUESTY MATCH
GROWN STATEMENT OF STATEME
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Search completed: June 7, 2003, 11:39:08 Job time: 89.3636 secs

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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 1000 summaries
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Maximum DB
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seq length: 25
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Match
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AI189662 qdl8h06.x
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D18747 MU3GS01809
AZ779522 2M0016K03
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AZ379234
7 BH75934781
7 AZ954781
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7 TA361F06P
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AE813511
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7 AZ335754
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AZ662093
AZ662609
TA186D08P
AZ405188
AZ427688
AZ427688
AZ548720
AZ56346716
AZ546716
   176129
AZ370656
AZ807609
AZ782243
TA359F10Q
AZ610564
TA23D03P
AZ3003P
AZ300687
AZ442914
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AZ589990
                                   AZG46716 1M0512D23
AZZ41B103 1M0512D23
AZZ41B103 1M0194103
AZZ25689 2M0107C22
AZZ399234 1M0134P16
BH759341 KG02510-3
AZZ94781 ZM022OC07
AZ435809 1M0223L03
AZS94781 ZM022OC05
AZ394781 ZM022OC05
AZ186405 GQ120005
AZ 1186405 GQ120005
AZ 1186405 GQ120005
AZ 1267790 T M00117A08
AL465023 T brucei
AL465023 T brucei
AZ 1267001 1M0361N23
AQ 17320 EC 1100384C12
AZ 173052 1M0584C12
AZ 173052 1M0584C11
AZ 186482 EC 11909
AZ 17256 AUZ57256
AZ 1725756 AUZ57260
AZ 1725756 AUZ57260
AZ 172579 1M0267801
AZ 172579 1M0267801
AZ 172579 1M054720
AZ 172672 1M054720
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AZ795704 2M0029C21
BHB13511 SALK 0643
AZ702000 1M0556P08
AL488403 T. bruce1
AA916047 0930d01.8
AZ592068 1M0402H08
AZ771198 1M0573M08
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AL477083 T. brucei
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39.0 23 24 39.0 23 25 39.0 23	153 8 40.0 25 154 8 40.0 25 155 8 40.0 25 157 8 40.0 25 158 8 40.0 25 159 8 40.0 25 161 8 40.0 25 162 8 40.0 25 163 7 8 39.0 19 165 7.8 39.0 19 167 7.8 39.0 20 169 7.8 39.0 20 170 7.8 39.0 20 171 7.8 39.0 20 172 7.8 39.0 20 173 7.8 39.0 20 174 7.8 39.0 20 175 7.8 39.0 20 177 7.8 39.0 20 178 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 20 179 39.0 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170 39.0 20 170 39.0 20 170 39.0 20 170 39.0 20 170 39.0 20 170 39.0 20 170 39
213     7.8     39.0     25       214     7.8     39.0     25       215     7.8     39.0     25       216     7.8     39.0     25       217     7.8     39.0     25       218     7.8     39.0     25       219     7.8     39.0     25       220     7.8     39.0     25       221     7.8     39.0     25       222     7.8     39.0     25       223     7.8     39.0     25       224     7.8     39.0     25       225     7.6     38.0     18	1193 7.8 39.0 23 1 1194 7.8 39.0 23 1 1195 7.8 39.0 23 1 1196 7.8 39.0 23 1 1197 7.8 39.0 23 1 1198 7.8 39.0 23 1 1298 7.8 39.0 24 1 1200 7.8 39.0 24 1 1201 7.8 39.0 24 1 1202 7.8 39.0 24 1 1203 7.8 39.0 24 1 1206 7.8 39.0 24 1 1207 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1	1177 7.8 39.0 21 17 179 7.8 39.0 21 17 180 7.8 39.0 22 9 1180 7.8 39.0 22 9 1181 7.8 39.0 22 11 1183 7.8 39.0 22 11 1184 7.8 39.0 22 11 1185 7.8 39.0 22 11 1186 7.8 39.0 22 11 1187 7.8 39.0 22 11 1187 7.8 39.0 22 11 1189 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11	153  8 40.0 25 9  154  8 40.0 25 9  155  8 40.0 25 17  158  8 40.0 25 17  159  8 40.0 25 17  161  8 40.0 25 17  162  8 40.0 25 17  163  8 40.0 25 17  164  7 8 40.0 25 17  165  7 8 39.0 19 17  168  7 8 39.0 20 17  169  170  170  171  183  190  171  171  183  190  171  171  173  178  189  190  170  171  171  171  171  171  17
213     7.8     39.0     25       214     7.8     39.0     25       215     7.8     39.0     25       216     7.8     39.0     25       217     7.8     39.0     25       218     7.8     39.0     25       219     7.8     39.0     25       220     7.8     39.0     25       221     7.8     39.0     25       222     7.8     39.0     25       223     7.8     39.0     25       224     7.8     39.0     25       225     7.6     38.0     18	1193 7.8 39.0 23 1 1194 7.8 39.0 23 1 1195 7.8 39.0 23 1 1196 7.8 39.0 23 1 1197 7.8 39.0 23 1 1198 7.8 39.0 23 1 1298 7.8 39.0 24 1 1200 7.8 39.0 24 1 1201 7.8 39.0 24 1 1202 7.8 39.0 24 1 1203 7.8 39.0 24 1 1206 7.8 39.0 24 1 1207 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1	1177 7.8 39.0 21 17 179 7.8 39.0 21 17 180 7.8 39.0 22 9 1180 7.8 39.0 22 9 1181 7.8 39.0 22 11 1183 7.8 39.0 22 11 1184 7.8 39.0 22 11 1185 7.8 39.0 22 11 1186 7.8 39.0 22 11 1187 7.8 39.0 22 11 1187 7.8 39.0 22 11 1189 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11	153  8 40.0 25 9  154  8 40.0 25 9  155  8 40.0 25 17  158  8 40.0 25 17  159  8 40.0 25 17  161  8 40.0 25 17  162  8 40.0 25 17  163  8 40.0 25 17  164  7 8 40.0 25 17  165  7 8 39.0 19 17  168  7 8 39.0 20 17  169  170  170  171  183  190  171  171  183  190  171  171  173  178  189  190  170  171  171  171  171  171  17
213     7.8     39.0     25       214     7.8     39.0     25       215     7.8     39.0     25       216     7.8     39.0     25       217     7.8     39.0     25       218     7.8     39.0     25       219     7.8     39.0     25       220     7.8     39.0     25       221     7.8     39.0     25       222     7.8     39.0     25       223     7.8     39.0     25       224     7.8     39.0     25       225     7.6     38.0     18	1193 7.8 39.0 23 1 1194 7.8 39.0 23 1 1195 7.8 39.0 23 1 1196 7.8 39.0 23 1 1197 7.8 39.0 23 1 1198 7.8 39.0 23 1 1298 7.8 39.0 24 1 1200 7.8 39.0 24 1 1201 7.8 39.0 24 1 1202 7.8 39.0 24 1 1203 7.8 39.0 24 1 1206 7.8 39.0 24 1 1207 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1	1177 7.8 39.0 21 17 179 7.8 39.0 21 17 180 7.8 39.0 22 9 1180 7.8 39.0 22 9 1181 7.8 39.0 22 11 1183 7.8 39.0 22 11 1184 7.8 39.0 22 11 1185 7.8 39.0 22 11 1186 7.8 39.0 22 11 1187 7.8 39.0 22 11 1187 7.8 39.0 22 11 1189 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11	153  8 40.0 25 9  154  8 40.0 25 9  155  8 40.0 25 17  158  8 40.0 25 17  159  8 40.0 25 17  161  8 40.0 25 17  162  8 40.0 25 17  163  8 40.0 25 17  164  7 8 40.0 25 17  165  7 8 40.0 25 17  167  7 8 39.0 19 17  168  7 8 39.0 20 17  169  170  171  171  183  190  171  171  171  171  171  171  171
213     7.8     39.0     25       214     7.8     39.0     25       215     7.8     39.0     25       216     7.8     39.0     25       217     7.8     39.0     25       218     7.8     39.0     25       219     7.8     39.0     25       220     7.8     39.0     25       221     7.8     39.0     25       222     7.8     39.0     25       223     7.8     39.0     25       224     7.8     39.0     25       225     7.6     38.0     18	1193 7.8 39.0 23 1 1194 7.8 39.0 23 1 1195 7.8 39.0 23 1 1196 7.8 39.0 23 1 1197 7.8 39.0 23 1 1198 7.8 39.0 23 1 1298 7.8 39.0 24 1 1200 7.8 39.0 24 1 1201 7.8 39.0 24 1 1202 7.8 39.0 24 1 1203 7.8 39.0 24 1 1206 7.8 39.0 24 1 1207 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1	1177 7.8 39.0 21 17 179 7.8 39.0 21 17 180 7.8 39.0 22 9 1180 7.8 39.0 22 9 1181 7.8 39.0 22 11 1183 7.8 39.0 22 11 1184 7.8 39.0 22 11 1185 7.8 39.0 22 11 1186 7.8 39.0 22 11 1187 7.8 39.0 22 11 1187 7.8 39.0 22 11 1189 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1191 7.8 39.0 23 11 1192 7.8 39.0 23 11	153  8 40.0 25 9  155  8 40.0 25 9  156  8 40.0 25 17  158  8 40.0 25 17  158  8 40.0 25 17  161  8 40.0 25 17  162  8 40.0 25 17  163  8 40.0 25 17  164  7 8 40.0 25 17  165  7 8 39.0 19 17  168  7 8 39.0 20 17  168  7 8 39.0 20 17  169  170  171  171  183  190  171  171  171  171  171  171  171
213     7.8     39.0     25       214     7.8     39.0     25       215     7.8     39.0     25       216     7.8     39.0     25       217     7.8     39.0     25       218     7.8     39.0     25       219     7.8     39.0     25       220     7.8     39.0     25       221     7.8     39.0     25       222     7.8     39.0     25       223     7.8     39.0     25       224     7.8     39.0     25       225     7.6     38.0     18	1193 7.8 39.0 23 1 1194 7.8 39.0 23 1 1195 7.8 39.0 23 1 1196 7.8 39.0 23 1 1197 7.8 39.0 23 1 1198 7.8 39.0 23 1 1298 7.8 39.0 24 1 1200 7.8 39.0 24 1 1201 7.8 39.0 24 1 1202 7.8 39.0 24 1 1203 7.8 39.0 24 1 1206 7.8 39.0 24 1 1207 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1 1208 7.8 39.0 24 1	1177 7.8 39.0 21 17 179 7.8 39.0 21 17 180 7.8 39.0 22 9 1180 7.8 39.0 22 9 1181 7.8 39.0 22 11 1183 7.8 39.0 22 11 1184 7.8 39.0 22 11 1185 7.8 39.0 22 11 1186 7.8 39.0 22 11 1187 7.8 39.0 22 11 1187 7.8 39.0 22 11 1189 7.8 39.0 23 11 1190 7.8 39.0 23 11 1190 7.8 39.0 23 11 1191 7.8 39.0 23 11 1192 7.8 39.0 23 11	153  8 40.0 25 9  155  8 40.0 25 9  156  8 40.0 25 17  158  8 40.0 25 17  158  8 40.0 25 17  161  8 40.0 25 17  162  8 40.0 25 17  163  8 40.0 25 17  164  7 8 40.0 25 17  165  7 8 39.0 19 17  168  7 8 39.0 20 17  168  7 8 39.0 20 17  169  170  171  171  183  190  171  171  171  171  171  171  171
213 7.8 39.0 25 9 AUZ54196 214 7.8 39.0 25 9 AVZ46508 215 7.8 39.0 25 10 AW246508 216 7.8 39.0 25 17 AZ394252 217 7.8 39.0 25 17 AZ483610 218 7.8 39.0 25 17 AZ7483610 219 7.8 39.0 25 17 AZ768392 220 7.8 39.0 25 17 AZ768392 221 7.8 39.0 25 17 AZ76699 222 7.8 39.0 25 17 BH789461 223 7.8 39.0 25 17 BH789461 224 7.8 39.0 25 17 BH789461 225 7.6 38.0 18 10 AW246520	193 7.8 39.0 23 17 AZ806458 194 7.8 39.0 23 17 AZ806458 195 7.8 39.0 23 17 AZ859707 196 7.8 39.0 23 17 AZ859702 197 7.8 39.0 23 17 TA242D06P 198 7.8 39.0 23 17 TA242D06P 199 7.8 39.0 24 17 TA39P09P 199 7.8 39.0 24 14 L32034 201 7.8 39.0 24 17 AZ341147 202 7.8 39.0 24 17 AZ340147 203 7.8 39.0 24 17 AZ3609979 204 7.8 39.0 24 17 AZ795736 205 7.8 39.0 24 17 AZ813287 206 7.8 39.0 24 17 AZ813287 207 7.8 39.0 24 17 AZ813287 208 7.8 39.0 24 17 AZ813280 209 7.8 39.0 24 17 AZ8132BD04Q 211 7.8 39.0 24 17 TA31BD04Q 212 7.8 39.0 24 17 TA5F09P 212 7.8 39.0 24 17 TA5F09P	177 7.8 39.0 21 17 AZ486594 178 39.0 21 17 AZ784525 179 7.8 39.0 21 17 AZ786205 180 7.8 39.0 22 9 AZ935965 181 7.8 39.0 22 9 AZ935965 182 7.8 39.0 22 17 AZ360276 183 7.8 39.0 22 17 AZ588797 184 7.8 39.0 22 17 AZ588797 185 7.8 39.0 22 17 AZ588707 186 7.8 39.0 22 17 AZ623402 187 7.8 39.0 22 17 AZ623402 188 7.8 39.0 22 17 AZ623402 189 7.8 39.0 22 17 AZ623402 189 7.8 39.0 22 17 AZ63707 189 7.8 39.0 22 17 AZ63707 189 7.8 39.0 22 17 AZ63707 189 7.8 39.0 22 17 AZ663707 189 7.8 39.0 23 17 AZ387861 190 7.8 39.0 23 17 AZ387863 190 7.8 39.0 23 17 AZ387863 191 7.8 39.0 23 17 AZ387863 192 7.8 39.0 23 17 AZ387863	153 154 155 156 157 158 158 160 158 158 160 158 160 159 161 161 162 163 164 178 165 165 165 166 178 189 199 199 199 199 199 199 199 199 19
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Length 20;

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EST 28-OCT-1998

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AU256704/c
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                                                                                                                                                                                                                                                                                       Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Japan
Tel: 81-743-72-5589
                                                                    Generation of expressed sequence tags from mouse brain Unpublished (2002)
                                                                                     Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Rodentia;
1 (bages 1 to 20)
Kato,K. and Matoba,R.
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                                                                                                                                       house mouse.
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Sciurognathi; Muridae;
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s musculus cDNA clone
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Eukaryota, Metazoa, C
Mammalia, Eutheria, F
1 (bases 1 to 25)
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                  13,
                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (infco@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 812 Std Error: 0.00
Seq primer: -40UP from Gibco
High quality sequence stop: 1
Location/Qualifiers
3 TGCGTTTGCCACTCAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                  25 bp mRNA linear EST 28-OCT-
qd18h06.x1 Soares placenta 8to9weeks 2NbHP8to9W Homo sapiens cf.
clone IMAGE:1724123 3' similar to TR:Q14211 Q14211 E4BP4 GENE.
; mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                   Tumor Gene Index 
Unpublished (1997)
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                           Similarity
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National Cancer Institute, Cancer Genome Anat
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nilarity 77.8%;
Conservative
                 Conservative
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                                                              double-stranded cDNA was size selected, ligated to Eco adapters (Pharmacia), digested with Not I and cloned the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library constructed by Bento Soares and M.Fatima Bonaldo."
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/db_xref="taxon:10000"
/clone_"BED0008872"
/clone_1b="3'-directed mouse cDNA library"
/tissue_type="brain"
/note="Vector: pGEM-T-easy"
3 c 6 g 3 t
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                        56.0%;
81.2%;
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Pred. No. 9.1e+04;
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AU257541 3'-directed mouse
BED0010829 3', mRNA sequen
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Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Japan
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1 (bases 1 to 24)
Kato, K. and Matoba, R.
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Mammalia; Eutheria;
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                                                                                                                        CDNA sequencing 
Unpublished (1995)
                                                                                                                                               Kawamoto, S., Okubo, K., YOSHIL
Analysis of gene expression
                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                     Mus musculus domesticus
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                                                                                                                                                                                                                                                                                       D18747.1
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URL:http://love2.aist-nara.ac.jp/BED/index.html.
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                                             Osaka University
3-1 Yamada-oka, Suita,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           touse mouse.
                                                                                                         Contact: Kawamoto, S.,
                                                                                                                                                                                                                                                    western European house mouse.
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                                                                         Institute for Cellular and Molecular Biology
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/note="Vector: pGEM-T-easy"
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/db_xref="taxon:10090"
organism="Mus musculus"
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                                                                                                                                                                                                                                                                                     GI:1100716
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clone_lib="3'-directed mouse cDNA library"
                                                                                                                                                        Okubo,K., Yoshii,J., Katsuki,M. and Matsubara,K. gene expression in mouse embryogenesis by 3'-directed
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73.7%;
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Rodentia;
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Pred. No. 1.9e+05
0; Mismatches
                                                                                                         Okubo, K., Yoshii, J., Katsuki, M. and
                                           Osaka 565, Japan.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0016 row: K column: 03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn, D., Aoyagi, A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Plate: 0016 row: K column: 03
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
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University of Utah
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pubp42 (gil4732114]gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F./note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackt Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical
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/db xref="taxon:10092"
/clone="md0270"
/clone_ib="Mouse 3'-directed"
/clone_type="decidual tissue (day 6.5-8.5 of
/tissue_type="decidual tissue (day 6.5-8.5 of
                                                                                                                                                                                                                                                                                                                               clone lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                             db_xref="taxon:10090"
clone="UUGC2M0016K03"
                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Mus musculus"
strain="C57BL/6J"
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Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ625339 19 bp DNA linear GSS 13-DEC-200100464F13R Mouse 10kb plasmid UUGCIM library Mus musculus genomic Clone UUGC1M0464F13 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi [473214 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated
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                                                                                                                                                                                                                                                                               /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                            clone="UUGC1M0464F13"
clone_lib="Mouse 10kb plasmid UUGC1M library"
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AU254819 3'-directed mouse cDNA library Mus musculus
BED0003462 3', mRNA sequence.
AU254819
AU254819 1 GI:20316963
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2002)
Contact: Kikuya Kato
Graduate School of Biological Sciences
Graduate School of Science and Technology
Nara Institute of Science and Technology
Nara Institute of Science and Technology
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Tel: 81-743-72-5581
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Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: kkato@bs.aist-nara.ac.jp/
URL:http://love2.aist-nara.ac.jp/BED/index.html
Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                     2M0029C21R Mouse 101
clone UUGC2M0029C21
AZ785704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fax: 81-743-72-5589
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                      /clone="BED0003462"
(clone lib="3'-directed mouse
/tissue_type="brain"
/note="Vector: pGEM-T-easy"
/note="3 c 2 t
                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
/db_xref="taxon:10090"
                                                                         GI:12922730
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10kb plasmid UUGCIM library
C21 R, DNA sequence.
   Chordata;
Rodentia;
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                                              SOURCE
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Best Local
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                                                                                                                         24 bp DNA linear GSS 02-MAY-201 SALK 064374 Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_064374, DNA sequence.

BH813511
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University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0029 row: C column: 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
Arabidopsis thaliana
Eukaryota, Viridiplantae, Streptophyta, Embryophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core et
                                                                   thale cress.
                                                                                                          BH813511.1 GI:20392116
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4 c 7 g 6 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC2M0029C21"
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/strain="C57BL/6J"
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Pred. No. 5.8e+05;
0; Mismatches 5;
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GSS.
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                                                                                                          USA
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Unpublished (2000)
Contact: Robert B. We University of Utah Go University of Utah Rm. 308, Biomedical I
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis Genome 
Unpublished (2001)
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A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0556 row: F column: 08
                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ762000 24 bp DNA linear GSS 16 1M0556F08R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC1M0556F08 R, DNA sequence.
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                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                     Mouse whole genome
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/clone="SALK_064374"
/clone_lib="Ārabidopsis thaliana TDNA insertion lines"
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TA259G08Q/c
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Best Local :
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T. brucei sheared genomic DNA
Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear trypanosoma tight size distribution ( 4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                   Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 15A, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                        Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Location/Qualifiers
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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A clone 259g08, reverse sequence;
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AA916047/c
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Matches 12; Conserv
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Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
unknown library type
unknown library type
std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersh
High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre
at http://www.sanger.ac.uk/Projects/T_brucei/.
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Og30d01.s1 NCI CGAP Br7 Homo sapiens CDNA clone IMAGE:1441345 3'
similar to SW:RL34_HUMAN P49207 60S RIBOSOMAL PROTEIN L34. ;, mRNA
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1_ (bases 1 to 22)
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National Cancer Institute, Cancer Genome Ana
AZ592068 22 bp DNA linear GSS 13-DEC-200
1M0402H08R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0402H08 R, DNA sequence.
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/strain="TREU927"
/db_xref="taxon:5691"
/clone="259g08"
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                                                                                                                                                                                                                                                             /note="Organ: breast; Vector: pCMV-SPORT4; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT Average insert size 1.2 kb. Life Technologies catalog #:10985-018" 7 g 5 t
                                                                                                                                                                                                                                                                                                                                                /clone_lib="NCI_CGAP_Br7"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="IMAGE:1441345"
                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Homo sapiens"
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84.6%;
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Pred. No. 9.3e+05;
0; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0402 row: H column: 08
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hanil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
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AZ771198 22 bp DNA li
1M0573M08F Mouse 10kb plasmid UUGC1M library
clone UUGC1M0573M08 F, DNA sequence.
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Fax: 801 585 7177
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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/clone="UUGC1M0402H08"
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/strain="C57BL/6J"
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pred. No. 9.3e+05;
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TA223E07P 22 bp. D
T. brucei sheared genomic DNA clone
genomic survey sequence.
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Insert Length: 10000 Std Erro
Plate: 0573 row: M column: 01
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coil XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0573M08"
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Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to give a tight size distribution (
4.kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T. brucei sheared genomic genomic survey sequence. AL477083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
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Melville, S.E., Rajandream, M.A. and Barrell, B.
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                                                                Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                     Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                          Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Ho Melville, S.E., Rajandream, M.A. and Barrell, B.G.
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                                                                                                                                    nh1@sanger.ac.uk
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1,N., Bowman,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L.,
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|strain="TREU927"
|db_xref="rexon:5691"
|clone="223e07"
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Pred. No. 9.3e+05;
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Email: nelsayedætigr.org
Details of T. brucei sequencing at the Sanger
at http://www.sanger.ac.uk/projects/T_brucei/.
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BH864166
                                                                                                                                                                                                                                                                                                                                                                                     Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                       Tel: 858 453 4100 x1752 Fax: 858 558 6379
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                                                                                                                                                                                                                                                                                                                                          Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2001)
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                                             /organism="Trypanosoma
/strain="TREU927"
/db_xref="taxon:5691"
/clone="202h11"
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Pred. No. 9.7e+05;
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RESULT 19
AZ589990/c
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SALK 095576 Arabidopsis thaliana TDNA insertion lines Arabidopsis
thaliana genomic clone SALK_095576, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                           12;
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                             AZ589990 20 bp DNA linear GSS 13-DEC-20 1M0399016F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0399016 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Pax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Zimmerman, J. and Ecker, J.R.

A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSS
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Unpublished (2001)
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                                                                                                                               AZ589990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="SALK_095576"
/clone lib="Arabidopsis thaliana TNNA insertion lines"
/note="PCFR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 3 c 3 g 13 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                           48.0%;
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Pred. No. 9.7e+05;
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Prednis, L.,
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RESULT 20
AZ812038/c
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Matches 1
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                                                                                                                                                                                                                                                                                                        14
                                                                                                                                                                                                                                                                                                                                                 10 GCCACTCAGAG 20
Mus musculus
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                            21 bp DNA linear GSS 20-FBB-20-
2M0078022F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0078022 F, DNA sequence.
AZ812038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0399 row: O column: 10
                                                                                                             AZ812038.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 20.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Utah Genome University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
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                                                                       house mouse.
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                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                        GCAACTCAGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Industrial PMP42Dry; Putified genomic DNA from M.
Industrial CS7BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0,005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWn42 (gil4732114/gb/AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coil XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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Lab_host="E. Coli strain XL10-Gold, T1-resistant, F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'clone_lib="Mouse 10kb plasmid UUGC1M library"
'sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      db_xref="taxon:10090"
clone="UUGC1M0399016"
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                                                                                                             GI:12980889
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      Chordata;
Rodentia;
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Pred. No. 1.1e+06;
0; Mismatches 1;
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    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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RESULT 21
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Best Local Similarity
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. and Wright, D., Weiss, R. Mouse, whole genome scaffolding with paired end reads from 10kb
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Unpublished (2000)
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University of Utah Genome Center
University of Utah
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutherla; Rodentia; Sciurognathi; Muridae; Nurinae; Mus
                                                                                                                                                                  AZ662093 22 bp DNA linear GSS 14-DEC-200
1M0541C05F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0541C05 F, DNA sequence.
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Ingert Length: 10000 Std Error:
Plate: 0078 row: O column: 22
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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                                                                                                                          AZ662093.1 GI:11799239
                                                                         house mouse.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWP42 (gil 4732114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab host="E. Coli strain XL10-Gold, Ti-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone="UUGC2M0078022"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/strain="C57BL/6J"
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REFERENCE
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Best Local Similarity
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
house mouse.
Mus musculus
Eukaryota, Metazoa; Chordata, Craniata, Vertebrata; Euteleostomi;
Euraryota, Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus
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Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0541 row: C column: 05
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                        1M0439P23F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0439P23 F, DNA sequence.
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Fax: 801 585 7177
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University of Utah
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                                                                                                                                                                      AZ612609.1 GI:11734715
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308,
USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWBA2 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XII0-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone_lib="Mouse 10kb plasmid UUGC1M library"
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clone="UUGC1M0541C05"
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strain="C57BL/6J"
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90.9%;
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and Wright,D., Welse,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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Location/Qualifiers
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Class: plasmid ends
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Insert Length: 10000 Std Erro
Plate: 0439 row: P column: 2
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                          T. brucei sheared genomic genomic survey sequence. AL475706
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
            Trypanosoma brucei.
Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                      AL475706.1 GI:11841017
GSS.
                                                                                                                                                                   TA186D08P
Тгураповота.
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Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="UUGC1M0439P23"
/clone_11b="Mouse 10kb plasmid UUGC1M library"
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90.9%;
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Pred. No. 1.2e+06;
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e 186d08,
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1 (bases 1 to 24)
Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R.,
Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
Melville, S.E., Rajandream, M.A. and Barrell, B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
Submitted (10-DEC-2000) Trypanosoma brucei genome Campus, Hinxton,
Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
                                                                      Insert Length: 10000 Std Error: Plate: 0173 row: G column: 15 Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutereoscom; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (Dases 1 to 21)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
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                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                             and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                              University of Utah
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/strain="TREU927"
/db_xref="taxon:5691"
/clone="186d08"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 21)
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                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0209 row: J column: 15
Seq primer: CACACAGGAAACAGCTATGACC
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quality sequence stop: 21.
Location/Qualifiers
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801 585 7177
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone lib="Mouse 10kb plasmid UUGC1M library"
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|strain="C57BL/6J"
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Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.
Plate: 0211 row: D column: 23
Seq primer: CACACAGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0211D23R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0211D23 R, DNA sequence.
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Fax: 801 585 7177
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/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/db_xref="taxon:10090"
/clone="UUGC1M0209J15"
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                                                                                                                                                                                                                                   Contact: Robert B. W
University of Utah G
University of Utah
Rm. 308, Biomedical
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and wright,D.,Welse,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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                                          Tel: 801 585 5606
Pax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0378 row: E column: 16
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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/note="Vector: PWD42nv; Purified genomic DNA from M.
/nusculus C57BL/6J (Temale) was obtained from the Jackson
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/strain="C57BL/6J"
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/clone="UUGC2M0211D23"
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Pred. No. 1.5e+06;
0; Mismatches 3;
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University of Utah Ge
University of Utah
Rm. 308, Biomedical F
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1M0490AllR Mouse 10kb plasmid UUGCIM
Clone UUGCIM0490All R, DNA sequence.
                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0490 row: A column: 11
Seq primer: CACACAGGAAACAGCTATGACC
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AZ634674.1 GI:11756864
                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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quality sequence stop: 23 
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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78.6%;
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weilse, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                     Tel: 801 585 5606
Fax: 801 585 7177
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length; 10000 Std Error: 0
Plate: 0512 row: D column: 23
Plate: 0512 row: D column: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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AZ646716.1 GI:11777462
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Location/Qualifiers
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/organism="Mus musculus"
/strain="C57BL/6J"
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78.6%;
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/db xref="Fuxxon:10090"
/clone="UUGC1M0512D23"
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/clone="UUGC1M0512D23"
/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
muscullus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resources documents/dnares/). The DNA
(http://www.jax.org/resources/documents/dnares/). The DNA
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adaptored vector DNA, and transformed into
chemically-competent E. coli XII0-Gold (Stratagene) cells
and selected for ampicillin resistance."

Y Match

46.0%; Score 9.2; DB 17; Length 23;
Local Similarity 78.6%; Pred. No. 1.5e+06;
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18103/c

SAZ418103

RAZ418103

INITION 1M0194L03F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0194L03 F, DNA sequence.

SSSION AZ418103

AZ418103.1 GI:10542116

ORDS GSS.

CE house mouse.

Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0194 row: L column: 03
Seg primer: CGTTCTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 24.
Location/Qualifiers Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 24) plasmid inserts Unpublished (2000) University of Utah University of Utah Mouse whole genome Contact: Robert B. Biomedical Weiss Genome Center scaffolding with paired end reads from 10kb Polymers Research Bldg., Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 0.00 20 <u>د</u> 2030 E., SLC,

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REFERENCE
AUTHORS
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AZ829689
LOCUS
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                                                                                                                                                                                                                                                                                                                                                                       Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 24)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                       plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2M0107C22F Mouse 10kb plasmid TUGC1M library Mus musculus genomic clone UUGC2M0107C22 F, DNA sequence.
                                                                                                                                                                                                                                                                 University of Utah Genome Center University of Utah
                                                                   Email: ddunn@genetics.utah.edu
Inmert Length: 10000 Std Error: 0.00
Plate: 0107 row: C column: 22
Seq primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           house mouse.
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                                                                                                                                                                                                                                          308,
                                                                                                                                                                                                                                                                                                                     cact: Robert B.
                                                 ss: plasmid ends
quality sequence stop: 24.
Location/Qualifiers
                                                                                                                                                                    801 585 5606
801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW042 (gil4732114 [gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                      USA
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                     Weiss
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Pred. No. 1.5e+06;
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JOURNAL COMMENT
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AUTHORS
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ORGANISM
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VERSION
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AZ379234/c
LOCUS
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                                                                                                                                                                                                                                  Contact: Robert B. We
University of Utah Ge
University of Utah
Rm. 308, Biomedical P
                                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.
                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0134 row: P column: 16
Seg primer: CGTTGTARAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                  Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                           plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 25)
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AZ379234.1
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quality sequence stop: 25
Location/Qualifiers
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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|strain="C57BL/6J"
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78.6%;
                                                                                                                                                                                                                                                                                   Genome Center
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Pred. No. 1.5e+06;
0; Mismatches 3;
                                                                                                                 0.00
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BASE COUNT
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AUTHORS
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ORGANISM
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BH759341
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VERSION
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Best Local S
Matches 11
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                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 25)
Levis,R., Hoskins,R., Liao,G., Mozden,N., Tsang,G., He,Y., Karpen,G., Bellen,H., Rubin,G. and Spradling,A.
The Berkeley Drosophila Genome Project Gene Disruption Project Unpublished (2001)
Contact: Gerald Rubin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera, Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BH759341 25 bp DNA linear GSS 12-MAR-2000 KG02510-3prime Drosophila melanogaster P(SUPOr-P) P element insertion lines Drosophila melanogaster genomic Sequence recovered from 3' end of P element, DNA sequence.
                                                                                                                                                   Email: gerry@fruitfly.berkeley.edu
Sequence recovery method was inverse PCR.
Sequence orientation is forward strand relative to 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BH759341
BH759341.1 GI:19352580
                                                                                                                                                                                                                             Berkeley Drosophila Genome Project
University of California, Berkeley
LSA Building, Berkeley, CA 94720-3200,
Pax: 5106439947
                                              The P element insertion position is base 1 in the 25 bases. This insertion position refers to the first base of the 8 base target recognition sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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transposon-tagged.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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78.6%;
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Pred. No. 1.5e+06;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                         source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Metazoa,
Mammalia, Eutheria,
1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ954781 19 bp ' DNA 1:
2M0220C07R Mouse 10kb plasmid UUGC2M library
clone UUGC2M0220C07 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0220 row: C Column: 07
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                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
High quality sequence stop: 19.
                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
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/db xref="taxon:7227"
/clone lib="Drosophila melanogaster P{SUPor-P} P element
insertion lines"
was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were
                                                                                                                              /lab host="B. coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (female) was obtained from the Jackson
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                                                                                                             Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0220C07"
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                                                                                    (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                               'sex="Female"
                                                                                                                                                                                                                       clone_lib="Mouse 10kb plasmid UUGC2M library"
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78.6%;
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Pred. No. 1.5e+06;
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AZ435809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error:
Plate: 022 row: L column: 03
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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(http://www.jax.org/resources/documents/dnares/). The DNI was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                              /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42ny; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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/clone="UUGC1M0223L03"
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/strain="C57BL/6J"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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2M0275004R Mouse 10kb plasmid UUGC2M library Mus musculus
clone UUGC2M0275004 R, DNA sequence.
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Fax: 801 585 7177
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University of Utah
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Insert Length: 10000 Std
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                          /lab_host="E. coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (female) was obtained from the Jackson
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                                                                                                                           Laboratory Mouse DNA Resource
                                                                                                                                                                                                                        /clone lib="Mouse 10kb plasmid UUGC2M library"
/sex="Female"
                                                                                                                                                                                                                                                                        'db_xref="taxon:10090"
'clone="UUGC2M0275004"
                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
|strain="C57BL/6J"
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17
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Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk

Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                         Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                           to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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Trypanosoma brucei
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IGCGTTTGC 9
                                        TGCGTTTGC 11
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                                                                                                                                                                                                                 /organism="Trypanosoma brucei"
/strain="TREU927"
/db_xref="texon:5691"
/clone="361f06"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared to give a tight size distribution ( 4 kb). The v+i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
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Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                           AI186405 22 bp mRNA linear EST 28-OCT-199 qd20e05.x1 Soares placenta 8to9weeks 2NbHP8to9W Homo sapiens cDNA clone IMAGE:1724288 3' similar to TR:015726 Q15726 MALIGNANT MELANOMA METASTASIS-SUPPRESSOR. /, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL ; contact the
                                                        Tumor Gene Index
Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7 TTTGCCACT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
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AI186405.1 GI:3737043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
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/strain="TREU927"
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AZ367707/c
                                                                         PEATURES
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                     University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ367707 22 bp DNA linear GSS 02-OCT-20 1M0117A08R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC1M0117A08 R, DNA sequence.
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High quality sequence stop: 1
                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Lengtch: 10000 Std Error: 0.00
Plate: 0117 row: A column: 08
Seq primer: CACACAGGAAACAGCTATGACC
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M.Fatima Bonaldo."
4 c 8 g 2 t
                                                                       Location/Qualifiers
organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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db_xref="taxon:9606"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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Pred. No. is the number of results predicted by chance to have a

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## SUMMARIES

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Method of identifying and quantifying s
Patent: WO 0196612A, 188 20-DEC-2001;
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Location/Qualifiers
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Stachybotrys chartarum
Eukaryota; Fungi; Ascomycota;
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Behr,M., Small,P., Schoolnik,G. and W.
Molecular differences between species
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Haugland, R. and Vesper, S.
Method of identifying and quantifying specific fungi and bacteria
Patent: US 6387652-A 106 14-MAY-2002;
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                                                                                                                                                                                                                                                                                               Myrothecium verrucaria. Myrothecium verrucaria
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                                                                                                                                                                                                                                Method of identifying and quantifying specific fungi
Patent: WO 0196612-A 106 20-DEC-2001;
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY (US)
                                                                                                                                                                                                                                                                                                                                    AX402622.1
                                                                                                                                                                                                                                                                Haugland, R. and Vesper, S.
                                                                                                                                                                                                                                                                                      Eukaryota; Fungi; Ascomycota; mitosporic Ascomycota; Myrothecium.
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Similarity 87.5%;
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                                                                                                                                 Similarity
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/db_xref="taxon:5532"
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Pred. No. 2.1e+05;
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Method for producing fusion proteins
Patent: Ep 0212532-A1 7 04-MAR-1987;
                                                                                                                                                                                                                                                                                               13;
                                                                                                                                                                                                                                                                                                                                                                                           Process for producing heme proteins Patent: US 5681725-A 2 28-OCT-1997; Location/Qualifiers
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                             Brinkmann, U. and Hoffmeyer, S.
Polymorphisms in human genes of cardiovascular regulators and their use in diagnostic and therapeutic applications
Patent: WO 0224747-A 189 28-MAR-2002;
Epidauros Biotechnologie AG (DE)
                                                                                                                                                    Sequence 189 from Patent AX404363 AX404363.1 GI:21437644
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Chan, H.W., Shel
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1 (bases 1 to 22)
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Pred. No. 3.1e+05;
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Sequence 190 from Patent
AX404364
                                                                                                   1 (bases 1 to 18)
Chan,H.W., Shelton,E.R., Baecker,P.A.,
and Barnett,J.W.
Method for producing fusion proteins
Patent: EP 0212532-A1 11 04-MAR-1987;
Location/Qualifiers
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                                                                                                                                                                               Unknown
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/note="artificial.sequence~n=a or
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/db_xref="taxon:32630"
/note="artificial sequence-n=t or
9 c 3 g 5 t
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Pred. No. 3.7e+05;
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Pred. No. 3.7e+05;
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l Similarity 85.
12; Conservative
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Tearev.S.A., Emerson,S.U. and Purcell,R.H.
Recombinant proteins of a Pakistani strain
use in diagnostic methods and vaccines
patent: US 6207416-A 88:27-MAR-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Single nucleotide polymorphisms in genes Patent: WO 0118250-A 1207 15-MAR-2001; WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEAU Pharmaceuticals, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 21)
Lander, E.S., Gargill, M., Ireland, J.S., Bolk, S., Daley, G.Q. and Mccarthy, J.J.
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AX096029
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/db_xref="taxon:9606"
6 c 10 g
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Other publication CA 2112028 940629
Other publication CN 1090325 940803
Other publication AU 5275993 940707
Other publication FI 935900 940629
Other publication BE 1007723 951010
Other publication BE 1007731 950516
Other publication BE 1006483 940913.
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1 (bases 1 to 25)
1 (bases 1 to 25)
Tsarev,S.A., Emerson,S.U. and Purcell,R.H.
Recombinant proteins of a Pakistani strain
use in diagnostic methods and vaccines
Patent: US 6287759-A. 88 11-SEP-2001,
Location/Qualifiers
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                    Sequence 17 from Patent EP0634490.
A42267
A42267.1 GI:2297757
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A38123
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/db_xref="taxon:32644"
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Other publication BR 9402834 950613
Other publication FR 9402839 950114
Other publication FI 943389 950116
Other publication CA 2128050 950116
Other publication NO 942652 950116
Other publication AU 6743294 950127
Other publication GB 2279955 950118
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De, B.E., Lahaya, A.,
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                                                                                                                                                                                                                                                                                   Expression system for novel pullulanase Patent: US 5736375-A 3 07-APR-1998; Location/Qualifiers
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Deweer, P. and Amory, A.
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/db_xref="taxon:32644"
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Deweer, P. and Amory, A.
Pullulanase producing microrganisms
Patent: US 5817498-A 3 06-OCT-1998;
Location/Qualifiers
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Sequence 3 from patent US
AR098234
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Deweer, P. and Amory, A.
Pullulanase, microorganisms which produce
preparation of this pullulanese and the us
Patent: US 6074854-A 3 13-JUN-2000;
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Similarity 86.7%;
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AR127034
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De Buyl,E., Lahaye,A.,
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Pred. No. 5.8e+05;
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Deweer, P. and Amory, A.
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Patent: US 5721127-A 3 24-FEB-1998;
Location/Qualifiers
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Pred. No. 5.8e+05;
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DEWER, P. and Amory, A.

Process for the saccharification of

Patent: US 5731174-A 3 24-MAR-1998,

Location/Qualifiers
                                               Peptide comprising hepatitis B surface antigen Patent: EP 0304578-A 35 01-MAR-1989; Epitec Limited
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Hepatitis B surface antigen vaccine Patent: US 6072049-A 25 06-JUN-2000; Location/Qualifiers
                                                                                                                                                                                                                Thoma, H.A.

Hepatitis B surface antigen vaccine Patent: US 6072049-A 26 06-JUN-2000;
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Kirschner, M.W., King, R.W. and Peters, J.-M.

Kirschner, M.W., King, R.W. and Peters, J.-M.

Kirschner, M.W., King, R.W. and Peters, J.-M.

Kasay and reagents for detecting inhibitors of degradation of cell cycle regulatory proteins patent: US 5726025-A 13 10-MAR-1998;

Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

1 (bases 1 to 25)
Fransen,L. and Devos,K.
NEW POLYPEPTIDES AND PEPTIDES, NUCLEIC ACIDS CODING FOR THEM, AND NEW POLYPEPTIDES AND PEPTIDES, NUCLEIC ACIDS CODING FOR THEM, AND THEIR USE IN THE FIELD OF TUMOR THERAPY, INFLAMMATION OR IMMUNOLOGY Patent: WO 9322437-A 43 11-NOV-1993;
INNOGENETICS NV (BE); FRANSEN LUCIA (BE)
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                                                                                         Sequence 43
AR085110
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Fransen,L., Devos,K., Van De Voorde,A. and Van Heuverswyn,H. Polypeptides and peptides, nucleic acids coding for them, and their
             Unclassified.
1 (bases 1 to 25)
Fransen, L., Devos, K.,
                                                     Unknown
                                                                  Unknown
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/db xref="taxon:9606"
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Pred. No. 5.5e+05;
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Location/Qualifiers
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AX322593
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AR021129
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Location/Qualifiers
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AX404362
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Polymorphisms in human genes of cardiovascular regulators and their use in diagnostic and therapeutic applications
Patent: WO 0224747-A 188 28-WAR-2002;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Sequence 188 from Patent
AX404362
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Polymorphisms in human genes of cardiovascular regulators and their use in diagnostic and therapeutic applications
Patent: WO 0224747-A 192 28-MAR-2002;
Epidauros Biotechnologie AG (DE)
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Sequence 192 from Patent
AX404366
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Podstad,O., Hovig,E., Engebraaten,O.,
CAPL-specific oligonucleotides and me
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Sequence 3 :
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Location/Qualifiers
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Fodstad,O., Hovig,E., Engebraaten.O., Maelandsmo,G. and a CAPL-specific oligonucleotides and methods of inhibiting
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Location/Qualifiers
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1 (bases 1) to 24)
1 (bases 1) to 24)
1 (o'Reilly,L., Puthalakath,H., Adams,J., O'Connor,L
1 (o'Reilly,L.) Puthalakath,H., Adams,J., O'Connor,L
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                                                                       ORGANISM
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                                                       EST.
Tetrahymena thermophila.
Tetrahymena thermophila
Tetrahymena thermophila
Tetrahymena, Clijophora; Oligohymenophorea;
Eukaryota; Alveolata; Ciliophora; Tetrahymena.
                                                                                                  BM394673 23 bp mRNA linear EST 1750072-2-5-D08.r.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
   920
Tel:
      Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606:
                                                                                               BM394673
                               ,J. and Klobutcher,L.
EST from Tetrahymena thermophila,
Unpublished (2002)
                                                1 (bases 1 to 23)
Turkewitz, A.P., Ka
                                                                                        BM394673.1
J E.
. 773
                                                                                         GI:18194726
                                                                                                                                                              Karrer, K.M.,
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AZ313630
AZ313630
AZ348213
AZ442098
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AZ473322
AZ475852
AZ484500
AZ486007
AZ48600772
AZ500772
AZ512414
                                                                                                                                            ALIGNMENTS
                                               Jahn, C.,
         IL 60637,
                                    strain CU428.1,
                                               Orias, E.,
         USA
                                               Kirk, K.E.,
                                    growing
                                                                                                               EST 17-JAN-2002
                                                                                                                                                                                                          391 2M0217P07
211 2M0218P15
308 2M0267D11
436 2M0281F18
535 2M0281H19
7 HUMGS01683
7 HUMGS01683
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1M0289H08
1M0294A18
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1M0313E17
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1M0234F12
1M0331H02
257PbH06
1M0370P10
1M0407E16
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2M0145M02
2M0164F06
2M0197L06
2M0212D23
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1M0436E13
1M0451M06
1M0454G13
1M0481C17
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1M0339N15
1M0357J21
1M0403M11
                                                                                                                                                                                                1M0030106
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2M0080P02
2M0105P13
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2M0130J11
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2M0115N20
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RESULT 3
AZ634834/c
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BM394673
LOCUS
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AZ634834 24 bp DNA linear GSS 13-DEC-2000 1M0490C19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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University of Chicago
920 E. 58th Street, Chicago,
Tel: 773 702 4374
Fax: 773 702 3172
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BM394673.1 GI:18194726
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Tetrahymena thermophila cDNA, mRNA sequence.
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Seq primer: T3
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: apturkew@midway.uchicago.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Turkewitz AP
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Turkewitz, A.P., Ka
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/strain="CU428.1"
/db xref="taxon:5911"
/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+, Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
a 11 c 8 g 3 t
                                                                                                                                                                                                                                                /db_xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/clone lib="Chilcoat processes on library
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
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                                                                                                                                                                                                                                                                                                                                                       strain="CU428.1"
                                                                                                                                                                                                                                                                                                                                                                      organism="Tetrahymena thermophila"
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81.2%;
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Pred. No. 6.2e+05;
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Pred. No. 3.
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VERSION
                                                   RESULT 4
AZ848503
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Matches 13; Conserv
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Insert Length: 10000 Std Exror: 0.00
Plate: 0490 row: C column: 19
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone UUGCIM0490C19 R, DNA sequence.
AZ634834
AZ634834.1 GI:11757024
AZ848503 23 bp DNA 11Dear USS 41-FBF-400
2M0149H02R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia, Eutheria,
1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Pax: 801 585 7177
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                                                                                                                                                                                   TGCGCCCGGATCCAGGC 18
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                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector NNA and transformed into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                                                                                                                                                                                                             adaptored vector DNA, and transformed into chemically-competent E. coli Xil0-Gold (Stratagene) cells and selected for ampicillin resistance."
7 c 6 g 4 t
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'clone lib="Mouse 10kb plasmid UUGCIM library"
'sex="Male"
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strain="C57BL/6J"
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lone="UUGC1M0490C19"
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Pred. No. 6.2e+05;
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                               GSS 21-FEB-2001
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AZ855545
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
19 b
2M0159F09R Mouse 10kb plasmid
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Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
Plate: 0149 row: H column: 02
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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801 585 7177
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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strain="C57BL/6J"
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clone="UUGC2M0149H02"
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91.7%;
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Pred. No. 7.6e+05;
0; Mismatches 1;
   19 bp DNA linear GSS 21-FEB-20
mid UUGCIM library Mus musculus genomic
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                                            RESULT 6
AZ791884/c
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Best Local Similarity
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AZ855545
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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GSS.
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Insert Length: 10000 Std Error: 0.00
Plate: 0159 row: F column: 09
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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  AZ791884 21 t
2M0043A02F Mouse 10kb plasmid
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                   GCCCGGATCC 12
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                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil 4732114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWD42nv, Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       clone_lib="Mouse 10kb plasmid UUGC1M library"
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/clone="UUGC2M0159F09"
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    bp DNA linear
i UUGC1M library Mus n
      Mus musculus
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AZ591658/c
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AZ591658 20 bp DNA linear GSS 13 1M0401F19R Mouse 10kb plasmid UUGCIM library Mus musculus
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0043 row: A column: 02
Seq primer: CGTTGTAAAACGACGGCCAGT
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AZ791884
AZ791884.1 GI:12935238
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 21)
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Location/Qualifiers
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                                          CTGCGCCCGGATCCAGGC 18
                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNP042 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/db_xref="taxon:10090"
/clone="UUGC2M0043A02"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Pred. No. 1.1e+06;
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Insert Length: 10000 Std Error: 0.00
Plate: 0401 row: F column: 19
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Unpublished (2000)
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AZ305158 21 bp DNA linear GSS 29 1M0005K11R Mouse 10kb plasmid UUGC1M library Mus musculus
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801 585 7177
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                        http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH92 (gil4732114/gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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|strain="C57BL/6J"
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/clone="UUGC1M0401F19"
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Rodentia;
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Pred. No. 1.7e+06;
0; Mismatches 4;
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AZ305158
AZ305158.1 GI:10341896
AZ834643 21 bp DNA linear GSS 20-FEB-2001
2M0117N24F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
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Insert Length: 10000 Stdt Error: 0.00
Plate: 0005 row: K column: 11
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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/strain="C57BL/6J"
/db_xref="ruaxon:10090"
/clone="DUGC1M0005K11"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Pred. No. 1.7e+06;
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(2000)
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(Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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AZ487506 24 bp DNA linear GSS 05 1M0317P10F Mouse 10kb plasmid UUGC1M library Mus musculus
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AZ834643
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Insert Length: 10000 Std Error:
Plate: 0117 row: N column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Pax: 801 585 7177
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/clone="UUGC2M0117N24"
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/strain="C57BL/6J"
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75.0%;
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Pred. No. 1.7e+06;
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LOCUS TA118G04P 25 bp DNA linear DEFINITION T. brucei sheared genomic DNA clone 118g04, forward
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AZ487506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: plasmid ends
High quality sequence stop: 24.
Location/Qualifiers
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University of Utah Genome Center
University of Utah
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Insert Length: 10000 Etd Error: 0.00
Plate: 0317 row: P column: 10
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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/note="Vector: PWD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC1M0317P10"
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/strain="C57BL/6J"
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75.0%;
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutherla; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 24)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Trypanosoma brucei
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AL463737
                                 Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                           Unpublished (2000)
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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                                                     Contact: Robert B.
University of Utah
University of Utah
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                           GI:11757024
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78.6%;
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                                                                                   Genome Center
                                                                                                          We188
                                       Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 9.2; DB Pred. No. 2.5e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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2.5e+06;
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8
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                                         20 S. 2030 E.,
                                                                                                                                                                          reads from 10kb
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VERSION
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                                                                                                                                                                                                                                                 TITLE
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Best Local
                                                                                                                                                                                                                                                                                                                                                                           ORGANISM
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               Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 707 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                         Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 25)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NALIONAL CANCER INStitute, Cancer Genome Anatomy Project (CGAP);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA909237 25 bp mRNA linear EST 09-JUN-199 0108C04.s1 Soares NFL T GBC S1 Homo sapiens cDNA clone IMAGE:1522854 3' Similar to TR:002402 O02402 INSOLUBLE PROTEIN. ;,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ende
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA909237
AA909237.1 GI:3048642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA sequence.
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                                                                                                                                                                                               Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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quality sequence stop: 1.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constrant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW042 (gij4732114 [gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Length: 10000 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: PWD42nv; Purified genomic musculus C57BL/6J (male) was obtained fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
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/clone="UUGCIM0490C19"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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78.6%;
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Pred. No. 2.5e+06;
D; Mismatches 3;
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AZ785457
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VERSION
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11;
                                                                                                                                                                                                   Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00

Plate: 0029 row: A column: 24

Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 23)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                    High quality sequence stop: 23.
                                                                                                                                                                                                                                                                                                                                                         84112,
                                                                                                                                                                                                                                                                                                                                                                                                University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                            Biomedical
               /organism="Mus musculus"
/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC2M0029A24"
/clone_lb="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
'lab_host="E. Coli strain XL10-Gold, T1-resistant,
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clone_lib="Soares_NFL_T_GBC_S1"
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78.6%;
                                                                                                                                                                ion/Qualifiers
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Pred. No. 2.5e+06;
D; Mismatches 3
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
                                                                                                                                                                                                          Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends High quality sequence stop: 23.
                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0076 row: C column: 04
                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
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9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                        34112, USA
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x: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|473214|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
                                                                                                               organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                         location/Qualifiers
                        clone lib="Mouse 10kb plasmid UUGC1M library"
|sex="Male"
                                                                  db_xref="taxon:10090"
clone="UUGC1M0076C04"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                50.0%;
_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                          Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                        Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9; DB 17; LC...
No. 3.1e+06;
0;
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AZ827973
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Best Local
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                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0104 row: P column: 11
Seq primer: CACACGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ827973 23 bp DNA linear GSS 20-FEB-200
2M0104P11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0104P11 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 GCGCCCGGATCC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ827973.1
                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coil XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Biomedical Polymers Research Bldg., 20
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                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0104P11"
                                                                                                                                                                                                          Location/Qualifiers
                                                     clone lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GI:12997881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              48.9%;
host="E. Coli strain XL10-Gold, T1-resistant, F-"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 8.8; DB 17;
Pred. No. 3.8e+06;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                     Std Error: 0.00
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SOURCE

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RESULT 18
AI594892/c
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Matches
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Best Local Similarity
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                                                                                                                                                                                               This clone was previously sequenced on the 5' data is from the 3' end
Trace considered overall poor quality
Possible reversed clone: similarity on wrong in High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                   Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL:; contact the
TMAGE Consortium (info@image.llnl.gov) for further information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Marra M/WashU-NCI Mouse EST Project 1999
WashIngton University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
Tel: 314 286 1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Waterston, R. and Wilson, R.
The WashU-NCI Mouse EST Project 1999
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marra,M., Hillier,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,R., Ritter,B., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 25)
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25 bp mRNA linear EST 21-APR-1999
vel2c06.xl Soares mouse NbMH Mus musculus cDNA clone IMAGE:817930
37. similar to SW:SUR_RAT Q09429 SULFONYLUREA RECEPTOR. [1] ;, mRNA
                                                                                                                                                                                                                                                                                                                                              MGI:486210
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AI594892
AI594892.1 GI:4603940
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
/organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="reaxon:10090"
/clone="IMAGE:817930"
/clone_lib="Soares mouse NbMH"
                                                                                                                                                                    Location/Qualifiers
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1. No. 3.8e+06;
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BH852925/c
BASE COUNT
ORIGIN
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ORIGIN
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BH852925
BH852925.1
                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BH852925 25 bp DNA linear GSS SALK 075769.37.15.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK 075769.37.15.x,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 GCCCGGATCCAG 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                           Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                This is single pass sequence recovered from the left border of TDNA. This sequence lies within 300 bases of the 5' end of
                                                                                                                                                                                                                                                                                                                                                                                                Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             thale cress.
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                   directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html" s 6 c 9 g 5 t
                                                                                             /strain="Columbia 0"
/db xref="taxon:3702",
/db xref="taxon:3702",
/clone="8A1K_075769.37.15.x"
/clone="BA1K_075769.37.15.x"
/clone="BA1K_075769.37.15.x"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            provided by Dr. Minoru Ko, Wayne State Univ. Library constructed and normalized by Bento Soares and M.Fatima
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8 c
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/dev_stage="4 weeks"
/lab_host="DH10B"
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Pred. No. 3.8e+06;
0; Mismatches 2;
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AA954509/c
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                                                                           RESULT 21
BM395601/c
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Matches 11
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les 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OnBidO5.61 Soares NFL T GBC_S1 Homo sapiens cDNA clone IMAGE:1563081 3' Similar to TR:Q24035 Q24035 ENA POLYPE:TIDE. CONTAINS element MSR1 repetitive element ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IMAGE Consortium (info@image.llnl.gov) for Trace considered overall poor quality Insert Length: 754 Std Error: 0.00 Seq primer: -40m13 fwd. ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 GCGCCCGGATCC 14
BM395601 23 bp mRNA linear EST 1. 5009-0-1-C04.t.1 Chilcoat/Turkewitz cDNA (large fraction) Tetrahymena thermophila cDNA, mRNA sequence.
BM395601
                                                                                                                                                                                                  3 GCGCCCGGATCCAGG 17
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 19)
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EST.
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This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                 /note="Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687229, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="IMAGE:1563081"
/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
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73.3%;
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                                                                                                                                                                                                                                                        Score 8.6; DB 9;
Pred. No. 4.6e+06;
                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                             Length 19;
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                                                        EST 17-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AZ410218 23 bp DNA linear GSS 03-OCT-20: 1M0182F23F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0182F23 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tetrahymena thermophila.
Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                               Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,B., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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EST.
                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0182 row: F column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Chicago
920 E. 58th Street, Chicago,
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                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                           Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                 plasmid inserts .
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                             and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 23)
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s: plasmid ends
quality sequen
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/note="Vector: BlueScript2 SK.; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
Proc. Natl. Acad. Sci USA, 98: 8709-8713."
6 C 6 9 2 t
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/db_xref="taxon:5911"
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Pred. No. 4.6e+06;
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                                                                                                                                                                             Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0256 row: F column: 09
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                               Mouse
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                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
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2M0256F09R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0256F09 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              house mouse
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="UUGC1M0182F23"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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FEATURES
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Tetrahymena thermophila
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Turkewitz, A.P., Karre, J. and Klobutcher, L.
                                                                                                                                                                                                                                                                                                                                            Contact: Turkewitz AP
Molecular Genetics and Cell Biology
University of Chicago
920 E. 58th Street, Chicago, IL 606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BM397860 24 bp mRNA linear EST 17 5009-0-38-B08.t.1 Chilcoat/Turkewitz cDNA (large fraction)
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                                                                                                                                                                                                                                       Seq primer:
                                                                                                                                                                                                                                                                                                Tel: 773 702 4374
Fax: 773 702 3172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from Tetrahymena thermophila, strain CU428.1, growing cells
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                                                                                                                                                                                                                                                               apturkew@midway.uchicago.edu
/db xref="taxon:5911"
/clone lib="Chilcoat/Turkewitz cDNA (large fraction)"
/note="Vector: BlueScript2 SK+; Details on library
preparation can be found in Chilcoat and Turkewitz (2001)
                                                                                                                      organism="Tetrahymena thermophila"
/strain="CU428.1"
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/db_xref="taxon:10090"
/clone="UUGC2M0256F09"
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73.3%;
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T. brucel sheared genomic genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                      insert libraries for whole genome shotgun sequencing small genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                         Bequence.
                                                                                                                                                                                                                                                                                                                                                                                 Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.oc.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Trypanosoma brucei.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to give a tight size distribution ( 4\ kb ). The v + i method used for the library construction described in detail in Smith, H. and Venter, J.C. (Making a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 15A, B-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AL495180.1 GI:11871567
                                                                  HUMXP8D1B Human placenta
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lingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L.,
ille,S.E., Rajandream,M.A. and Barrell,B.G.
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/strain="TREU927"
/db xref="texon:5691"
/clone="368a07"
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AZ855545/c
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Isolation of chromosome-specific genes by reclaration of chromosome-specific genes by reclaration of cosmid libraries
                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0159 row: F column: 09
Seg primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                  Class: plasmid ends
                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                 University of Utah (
University of Utah 
Rm. 308, Biomedical
                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B.
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/clone="XP8D1B"
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               /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0159F09"
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/db_xref="taxon:9606"
clone_lib="Mouse 10kb plasmid UUGC1M library"
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73.3%;
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Pred. No. 4.7e+06;
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Lindsay,E.A., Zhao,Z.-Y.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12
                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0163 row: D column: 08
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 19.
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                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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                          organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0163D08"
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clone_lib="Mouse 10kb plasmid UUGC1M library"
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90.0%;
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RESULT 28 AZ858446 LOCUS

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                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah Genome Sessearch Bldg., 20 S. 2030 E., SLC,
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ858446 19 bp DNA linear GSS 21-FEB-200200163D08R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0163D08 R, DNA sequence.
                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 013 row: D column: 08
Seq primer: CACACAGCAACAGCTATGACC
Class: plasmid ends
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                quality sequence stop: 19.
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                                                                                                                                                                                                                                                                                                                                                                                             USA
/db xref="taxon:10090"
/clone="UUGC2M0163D08"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                             /organism="Mus musculus"
/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GI:13051622
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Pred. No. 5.7e+06;
0; Mismatches 1;
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weisse, R.
                                                                                                                                                                                                                                  Tel: 801 585 5606
Pax: 801 585 7177
Email: ddunn@genetics.utah.edu
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0476 row: A column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1M0476A13F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0476A13 F, DNA sequence.
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                                                                                                                                                                                  Class: plasmid ends
High quality sequence stop: 20.
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                      /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0476A13"
clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                           ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                          Std Error: 0.00
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                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0534 row: B column: 02
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah Genome Center 
University of Utah
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Unpublished (2000)
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Fax: 801 585 7177
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.gh quality sequence stop: 20.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                           USA
/organism="Mus musculus"
(strain="C57BL/G0"
/db_xref="traxon:10090"
/clone="UUGC1M0534B02"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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/note="Vector: PWD42nv, Purified genomic DNA from M.
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Best Local
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0229 row: P column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                   Class: plasmid ends
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
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1 (bases 1 to 20)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 bp DNA linear G3S 27-APR-20
2M0229P20F Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0229P20 F, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                USA
/db_xref="taxon:10090"
/clone="UUGC2M0229P20"
/clone_lib="Mouse 10kb plasmid UUGC2M library"
                                                                         organism="Mus musculus"
/strain="C57BL/6J"
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/note="Vector: PWD42nv, Purified genomic DNA from M.
/nusculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Laboratory Mouse DNA Resource
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1 (Dases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                           Trace considered overall poor quality
Insert Length: 760 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                     CDNA Library Preparation: M. Bento Soares, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                    www-bio.linl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1602940"
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                                                                                                            ocation/Qualifiers
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90.0%; Pred. No. 5.7e+06;
tive 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 812 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (Dases 1 to 22)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
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                                                                             /dev stage="20 week-post conception fetus"
/lab hoste "DH108 (ampicillin resistant)"
/note="Organ: Liver and Spleen; Vector: pT7T3D (Pharmacia)
/note="Organ: Liver and Spleen; Vector: pT7T3D (Pharmacia)
with a modified polylinker; Site 1: Pac I; Site 2: Eco RI;
This is a subtracted version of The original Soares fetal
liver spleen inFLS library. ist strand cDNA was primed
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                       clone_lib="Soares_fetal_liver_spleen_INFLS_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Homo sapiens"
db_xref="taxon:9606"
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46.7%;
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Pred. No. 5.7e+06;
0; Mismatches 1
Score 8.4; DB 9;
Pred. No. 5.7e+06;
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                                                                                                                                                                                                                                                           AII19986 25 bp mRNA linear uc25b01.rl Soares mammary_gland_NbMMG Mus musculus IMAGE:1398985 5' similar to TR:000399 000399 WS-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA878803
AA878803
25 bp mRNA linear EST 14-APR of 86e04.81 NCI CGAP Li5 Homo sapiens cDNA clone IMAGE:1437246 similar to SW:PRPM HUMAN P10161 SALIVARY PROLINE-RICH PROTEIN contains TAR1.t3 TAR1 repetitive element; mRNA sequence.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (baese 1 to 25)

Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T. Geisel,S., Kucaba,T., Lacy,M., Lan,, Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 25)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insert Length: 185 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
Email: cgapbs-romail.nih.gov
unknown library type
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index 
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo, sapiens
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                                                                                                                                      Mus musculus
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                                                                                                                                                                                                   AI119986.1
                                                                                                                                                             house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon: 9606"
/clone="IMAGE:1437246"
/clone_lib="NCI_CGAP_Li5"
/tissue_type="hepatic adenoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: liver; Vector: pCMV-SPORT4; Site_1: Sall; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 0.8 kb."
                                                                                                                                                                                                   GI:3520310
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Pred. No. 5.7e+06;
0; Mismatches 1
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REPERENCE
AUTHORS
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AI913416
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ORIGIN
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VERSION
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                                                                   COMMENT
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                                                                                                                      TITLE
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Best Local
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                                                                                   JOURNAL
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Possible reversed clone: similarity on wrong strand
Seq primer: -28ml3 rev2 ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                      A1913416 EST 16-DEC TE777a09.x1 NCI CGAP Pan1 Homo sapiens CDNA Clone IMAGE:2294584 similar to SW:PRPP HUMAN P81489 SALIVARY PROLINE-RICH PROTEIN CONTAINS PTR5.t2 MSR1 repetitive element; mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FMAX: 314 200 1010
Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WashIngton University School of MedicineP
WashIngton University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lc
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome
                                                                                                                                                          Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 25)
                                                                                                                NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Anav
                                                                                                                                                                                                                                  human
                                                                                                                                                                                                                                                                       AI913416.1 GI:5633271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Waterston, R.
The WashU-HHMI Mouse EST Project
                                                                                 Tumor Gene Index
Unpublished (1997)
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TGCGCCCGGA 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T 3'); double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. RNA provided by Dr. Minoru Ko, Wayne State Univ. Library constructed and normalized by Bento Soares and M.Fatima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia ) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bonaldo."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Soares_mammary_gland_NbMMG"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:10090"
/clone="IMAGE:1398985"
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|strain="C57BL/6J"
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dev_stage="4 weeks"
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Pred. No. 5.7e+06;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 25;
                                                                                                                    Anatomy Project (CGAP),
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       Sequencing
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AZ307545/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUTHORS
                                                                                                                    Bource
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                                                                                                                                                                                                                                                                                                                         308,
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Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0009 row: H column: 27
                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. University of Utah University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 25) Dunn D., Aoyagi.A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1M0009H22F Mouse 10kb plasmid UUGC1M library Mus clone UUGC1M0009H22 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Trace considered overall poor quality Insert Length: 1741 Std Error: 0.00 Seq primer: -40UP from Gibco
                                                                                                                                                                                        Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                           84112, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AZ307545.1 GI:10346652
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/organism="Mus musculus"
(strain="C578L/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0009H22"
/clone_1ib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                               Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Organ: pancreas; Vector: pCMV-SPORT6; Site 1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11548-013"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:2294584"
/clone_lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH108"
                                                                                                                                                ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                         Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                               Polymers Research Bldg.,
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Pred. No. 5.7e+06;
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Query Match
Best Local S
Matches 9
Query Match
Best Local Similarity 76.9
Matches 10; Conservative
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Tetrahymena thermophila.
Eukaryota, Alveolata, Ciliophora, Oligohymenophorea,
Hymenostomatida, Tetrahymenina, Tetrahymena.
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50072-2-11-B12.r.1 Chilcoat/Turkewitz cDNA (large fraction)
Tetrahymena thermophila cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Chicago
920 E. 58th Street, Chicago, IL 60637, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Turkewitz AP
Molecular Genetics and Cell Biology
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                                                                                                                                                                                                                                                                                                                                                                                                                            apturkew@midway.uchicago.edu
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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/clone_lib="Chilcoat/Turkewitz cDNA (large fraction)"
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/note="Vector: BlueScript2 SK+; Details on library
/note="Twetton on be found in Chilcoat and Turkewitz (2)
proc. Natl. Acad. Sci USA, 98: 8709-8713."
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76.9%;
Score 8.2; DB 13;
Pred. No. 6.9e+06;
0; Mismatches 3;
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RESULT 39 BM393804/c

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BASE COUNT

VERSION KEYWORDS

ACCESSION DEFINITION

SOURCE

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**FEATURES** 

BASE COUNT

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REFERENCE
AUTHORS
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BM394186/c
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Eukaryota; Alveolata; Ciliophora; Oligohymenophorea;
Hymenostomatida; Tetrahymenina; Tetrahymena.
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920 E. 58th Street, Chicago, IL 60637,
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Molecular Genetics and Cell Biology
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                                                                                                                                                              h 45.6%;
Similarity 76.9%;
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Fax: 773 702 3172
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/db xref="taxon:5911"
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preparation can be found in Chilcoat and Turkewitz (2001)
proc. Natl. Acad. Sci USA, 98: 8709-8713."
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Pred. No. 6.9e+06;
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; OTHER INFORMATION:
US-10-080-959A-5
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US-10-080-959A-5
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5, Application US/10080959A
Publication No. US20030054369A1
GENERAL INFORMATION:
APPLICANT: Cruz-Perez, Patricia
APPLICANT: Buttner, Mark P.
TITLE OF INVENTION: Method for Detection of Stachybotrys chartarum in Pure Culture
TITLE OF INVENTION: Samples Using Quantitative Polymerase Chain Reaction
FILE REFERENCE: 0001-00001
CURRENT APPLICATION NUMBER: US/10/080,959A
CURRENT TILING DATE: 2002-05-16
PRIOR APPLICATION NUMBER: US 60/280,712
PRIOR FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 5
SEQ ID NO 5
LENGTH: 18
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Best Local
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ORGANISM: Stachybotrys chartarum
FEATURE:
NAME/KEY:
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                                                                                              CTGCGCCCGGATCCAGGC 18
                                                                                                                                                                                    100.0%;
ilarity 100.0%;
Conservative (
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9 US-09-907-575-242

9 US-09-902-759-242

9 US-09-902-751-242

9 US-09-902-713-242

9 US-09-902-713-36

9 US-09-902-713-36

9 US-09-904-6-33-36

9 US-09-904-185-344

9 US-09-940-185-1514

10 US-09-940-185-1514

10 US-09-940-185-1585

10 US-09-940-185-2408

10 US-09-940-185-2835

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10 US-09-940-185-2835

10 US-09-940-185-2835

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10 US-09-940-185-2836

10 US-09-940-185-236

10 US-09-94
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Pred. No. 16;
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RESULT 4
US-10-241-258-28/c
US-10-241-258-28/c
; Sequence 28, Application US/10241258
; Publication No. US20030078206A1
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GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
APPLICANT: Lasser, Gerald W.
APPLICANT: Bishop, Paul D.
IITLE OF INVENTION: INHIBITORS FOR USE IN HEMOSTASIS AND IMMUNE FUNCTION FILE REFERENCE: 99-12C3
CURRENT FILING DATE: 2002-06-25
PRIOR APPLICATION NUMBER: 09/253,604
PRIOR APPLICATION NUMBER: 09/253,604
PRIOR FILING DATE: 1999-02-19
PRIOR APPLICATION NUMBER: 09/253,604
PRIOR FILING DATE: 1999-01-22
PRIOR FILING DATE: 1999-02-19
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 28
LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
COTTER INFORMATION: Oligonucleotide ZC15035
US-10-180-762-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/053,154
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 52
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 28
LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Oligonucleotide ZC15035
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US-10-180-762-28/c
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                                                                  Query Match
Best Local S
Matches 14
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Publication No. US20030022838A1
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APPLICANT: Sheppard, Paul O.
TITLE OF INVENTION: ANTIBODIES THAT BIND
TITLE OF INVENTION: ADJPOCYTE-SPECIFIC P
FILE REFERENCE: 97-30D1
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Best Local :
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CURRENT FILING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: 09/118,408
PRIOR FILING DATE: 1998-07-17
20 CCGAGCCCGGATCCATG 4
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Pred. No. 8e+03;
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                                                                  RESULT 6
US-09-423-800-34/c
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U8-10-241-258-28
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SOPTWARE: Microsoft Office 97
SEQ ID NO 54
LENGTH: 19
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ramos, Juan
APPLICANT: Sariaslani, Sima
APPLICANT: Sariaslani, Sima
TITLE OF INVENTION: Method for the Production of p-Hydroxybenzcate in Species of
TITLE OF INVENTION: Pseudomonas and Agrobacterium
FILE REFERENCE: BC1018 US CIP
CURRENT APPLICATION NUMBER: US/09/997,664
CURRENT FILING DATE: 2001-11-28
PRIOR APPLICATION NUMBER: 09/585,174
PRIOR FILING DATE: 2000-06-01
             Sequence 34, Application US/09423800 Patent No. US20020165363A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                        Query Match
Best Local 9
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LENGTH: 24
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OTHER INFORMATION: Description of Artificial Sequence: primer
OTHER INFORMATION: primer used for sequencing pcu
-09-997-664-54
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TITLE OF INVENTION: IMHIBITORS FOR USE IN HEMOSTASIS AND TITLE OF INVENTION: IMMUNE FUNCTION

PILE REFERENCE: 99-12
CURRENT APPLICATION NUMBER: US/10/241,258
CURRENT FILLING DATE: 2002-09-10
NUMBER OF SEQ ID NOS: 50
SOPTMARE: FastSEQ for Windows Version 3.0
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                        64.4%;
77.8%;
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Pred. No. 1.6e+04;
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; OTHER INFORMATION: Synthetic DNA
US-10-182-018-34
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US-10-182-018-34/c
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CURRENT APPLICATION NUMBER: US/09/423,800

CURRENT FILING DATE: 1999-11-12

PRIOR APPLICATION NUMBER: PCT/JP98/02116

PRIOR FILING DATE: 1998-05-13

PRIOR APPLICATION NUMBER: JP 125505/1997

PRIOR FILING DATE: 1997-05-15

PRIOR APPLICATION NUMBER: JP 194445/1997

PRIOR FILING DATE: 1997-07-18

NUMBER OF SEQ ID NOS: 87
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LENGTH: 19
                                                                                                     Sequence 496, Application US/10085906
Publication No. US20030054371A1
GENERAL INFORMATION:
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Publication No. US20030049211A1
GENERAL INFORMATION:
APPLICANT: CHUGAI SEIYAKU KABUSHIKI KAISHA
TITLE OP INVENTION: THERAPEUTIC AND PREVENTIVE AGENTS FOR DENTAL DISEASES
FILE REFERENCE: PH-1092-PCT
CURRENT APPLICATION UNMER: US/10/182,018
CURRENT FILING DATE: 2002-07-24
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Best Local Similarity
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 APPLICANT: Ying, Vincent
APPLICANT: Wu, Paul
APPLICANT: Gray, Gary S.
TITLE OF INVENTION: POLYMORPHIC ELEMENTS IN THE
TITLE OF INVENTION: COSTINULATORY RECEPTOR LOCUS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: JP 2000-83034
PRIOR FILING DATE: 2000-01-25
NUMBER OF SEQ ID NOS: 75
SOFTWARE: Patentin Ver. 2.0
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FILE REFERENCE: GNN-5343CP2
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TYPE: DNA
ORGANISM: Artificial Sequence
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92.3%;
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Pred. No. 2e+04;
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Pred. No. 2e+04;
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RESULT 9
US-09-990-385-20/c
; Sequence 20, Application US/09990385
; Publication No. US20020192771A1
; GENERAL INFORMATION:
; APPLICANT: Koji YANAI et al.
; APPLICANT: Koji YANAI et BETA-FRUCTOFURANOSIDASE AND ITS GENE, METHOD OF INVENTION: BETA-FRUCTOFURANOSIDASE GENE, SYSTEM FOR PRODUCI BETA-FRUCTOFURANOSIDASE, AND BETA-FRUCTOFURANOSIDASE VARIA
                                                              STRANDEDNESS: No. US20020192771A1 relevant
TOPOLOGY: Linear
MOLECULE TYPE: Synthetic DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 20:
US-09-990-385-20
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LENGTH: 21
Query Match 63.3%;
Best Local Similarity 92.3%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/126,215
PRIOR PILING DATE: 1999-03-25
PRIOR APPLICATION NUMBER: US 09/534,061
PRIOR PILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: PCT/US00/07938
PRIOR PILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 545
SOPTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/085,906
CURRENT FILING DATE: 2002-02-27
                                                                                                                                                                                                   TELEPHONE: 202-721-8250
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                          PILING DATE: 10-Sep-1998

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/142,623

FILING DATE: September 10, 1998

ATTORNEY/AGENT INFORMATION:

NAME: Lee Cheng

REGISTRATION NUMBER: 40,949

REFERENCE/DOCKET NUMBER: 2001-1611

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
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                                                                                                                                                                  TYPE: Nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/990,385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Wordperfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
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92.3%;
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Score 11.4; DB:
Pred. No. 2e+04;
0; Mismatches
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Pred. No. 2e
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                                 DB 9; Length 22;
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Indels
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JOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: US-09-961-077-531
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US-09-961-077-531
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                                    Matches
                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                     TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 531:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL
                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE DOCKET NUMBER: 219/247
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Merlo, Donald J.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR MODULATION OF GENE EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: September 2, 1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1263
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zwick, Michael G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16 GCCCGGATCCCGG 4
1 CTGCGCCCGGATCCAG 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION
                                                                                                                                               LENGTH: 16 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/679,645
FILING DATE: July 12, 1996
APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Comparible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/961,077
FILING DATE: 21-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09961077
. US20030014775A1
                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Skokut, Thomas ,
Young, Scott A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Guo, Lining
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edington, Brent E.
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                                                    68.8%;
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                                  Score 11.2; DB 9;
Pred. No. 2.6e+04;
2; Mismatches 3;
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                                                                     Length 16;
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CUGCGGCCUCAUCCAG

16

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RESULT 13
US-09-930-423-384/c
US-09-930-423-384/c
Sequence 384, Application US/09930423
Publication No. US20030092003A1
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APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REPERENCE: 400/022 (MBH800-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT PILLM DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOFTWARE: Patentin version 3.0
SEQ ID NO 2821
LENGTH. 17
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; OTHER INFORMATION: Description of Artificial Sequence:
US-09-864-785-2821
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US-09-864-785-2821
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US-09-864-785-335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3329
SOFTWARE: Patentin version 3.0
                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 335, Applic Patent No. US2002017 GENERAL INFORMATION:
                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PEATURE:
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                                                                                                                                                               Conservative
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Pred. No. 2.6e
2; Mismatches
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Pred. No. 2.
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US-09-866-108-651/c
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US-10-020-141-15
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                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.0 SEQ ID NO 384
                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                    Sequence 651,
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Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                         APPLICANT:
                                                                                                                                           APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggar
APPLICANT: PENN, Shar:
APPLICANT: HANZEL, Dav
                                                  APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/327,485
PRIOR FILING DATE: 2001-10-05
NUMBER OF SEQ ID NOS: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/020,141
CURRENT FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/313,097
PRIOR FILING DATE: 2001-08-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: McCarthy, Jeanette
APPLICANT: Ableson, Allen
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF VASCULAR DISEASE
FILE REFERENCE: MMI-002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for
FILE REFERENCE: MBHB00,918-A 400/027
CURRENT APPLICATION NUMBER: US/09/930,423
CURRENT FILING DATE: 2001-08-15
NUMBER OF SEQ ID NOS: 4553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ribozyme Pharmaceuticals,
                CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 17
TYPE: RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 17
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo Sapiens
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   APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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Similarity 81.2%;
                                                                                                 JI, Yonggang
PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
                                                                                                                                                                                                                                                                                                                                                                                  TGCGCCCGGATCCAGG 17
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Vo. US20030092013A1
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60/207,456
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Pred. No. 2
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APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04 APPLICATION NUMBER: US 60/236,359

2000-05-26

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US-09-866-108-652/c
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LENGTH: 17
TYPE: DNA
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US-09-866-108-651
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                                                                                                                        CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/866,108
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                                                                                                                                                                                                                                                           FILE REFERENCE: AEOMICA-
                                                                                                                                                                                                                                                                       PPLICANT: RANK, David R.
PPLICANT: CHEN, Wensheng
PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OR FILING DATE: 2001-01-30

RAPPLICATION NUMBER: PCT/US01/00662

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00661

OR FILING DATE: 2001-01-30

OR APPLICATION NUMBER: PCT/US01/00670

OR APPLICATION NUMBER: PCT/US01/00670

OR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2001-01-30
APPLICATION UNMER: PCT/US01/00668
FILING DATE: 2001-01-30
APPLICATION UNWEER: PCT/US01/00663
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
                                          APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
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FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
APPLICATION NUMBER: PCT/US01/00667 PILING DATE: 2001-01-30
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Similarity 81.2%;
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HANZEL, David K.
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Pred. No. 2
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US-09-961-077-621
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US-09-866-108-652
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SOFTWARE: Aecmica Sequence Listing Engine
SEQ ID NO 652
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 621, Appendication No.
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Best Local (
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PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00661
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CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/961,077
FILING DATE: 21-Sep-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30
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                                                                                                                                                ZIP: 90071-2066
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 60/234,687
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1263
                                                                                                                                                                                                                                                                                                                                                        TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                 STATE: California COUNTRY: U.S.A.
                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.
                                                                                                                                                                                                                  CITY: Los Angeles
                                                                                                                                                                                                                                                   STREET: 633 West Fifth Street
                                                                                                                                  MEDIUM TYPE: 3.5" Diakette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/09961077
o. US20030014775A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zwick, Michael G.
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Young, Scott A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Edington, Brent E.
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                                                                                                                  storage
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Pred. No. 2.
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                                                                                   DOS 5.0
                                                                                                                                  1.44 Mb
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RESULT 19
US-10-118-100-2/c
### Sequence 2, Application US/10118100
### Publication No. US20030077613A1
### Publication No. US20030077613A1
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US-09-734-188-37/c
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/ TYPE: DNA
/ ORGANISM: Homo sapiens
US-09-734-188-37
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goldman, Assistant Counsel, Robin A.
APPLICANT: New York State Office of Mental Health
APPLICANT: Nathan S. Kline Institute for Psychiatric Research
TITLE OP INVENTION: PAH
FILE REPERENCE: Kline Inst.
CURRENT APPLICATION NUMBER: US/09/734,188
CURRENT FILING DATE: 2000-12-12
                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 62
SOFTWARE: PatentIn Ver. 2.0
                                                   APPLICANT: BioInvent International AB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/253,025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Richardson Ph.D., Mary Ann
APPLICANT: Goldman, Assistant Counse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 621:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 219/247
TELECOMMUNICATION IMFORMATION:
TELEPHONE: (213) 489-1600
OF INVENTION: A Method For In Vitro Molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                      TGGGCTCGGATCCATG 2
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FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/679,645
                 Soderlind, Ul.
Borrebaeck, C.
                                                                                                                                                                                                                                                                           Conservative
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81.2%;
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Pred. No. 2.5e+04;
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Pred. No. 2.
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US-10-118-100-36/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-813-289-3
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CURRENT FILING DATE: 2002-04-08
PRIOR APPLICATION NUMBER: US/09/341,711
PRIOR FILING DATE: 1999-09-21
PRIOR APPLICATION NUMBER: PCT/GB98/00219
PRIOR FILING DATE: 1998-01-26
PRIOR APPLICATION NUMBER: GB9701425.2
PRIOR APPLICATION NUMBER: GB9701425.2
PRIOR FILING DATE: 1997-01-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Fa
SEQ ID NO 3
LENGTH: 22
                                                                                                                                                           Sequence 36, Application US/10118100 Publication No. US20030077613A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/09813289
Patent No. US20020061571A1
GENERAL INFORMATION:
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SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 2

LENGTH: 22

TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                          Matches
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APPLICANT: BioInvent International AB
APPLICANT: Soderlind, Ulf
APPLICANT: Soderlind, Ulf
APPLICANT: Borreback, Carl
TITLE OF INVENTION: A Method For In Vitro Molecular
TITLE OF INVENTION: Evolution of Protein Function
FILE REFERENCE: Mewburn 341711
CURRENT APPLICATION NUMBER: US/10/118,100
CURRENT FILING DATE: 2002-04-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: No. US20020061571A1el isoform of myotonic dystrophy associated prilitle OF INVENTION: thereof
FILE REFERENCE: 800.027US1
CURRENT APPLICATION NUMBER: US/09/813,289
CURRENT FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,590
PRIOR PILING DATE: 2000-03-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Mahadevan, M.S. APPLICANT: Tiscornia, G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Evolution
FILE REFERENCE: Mewburn 341711
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Pred. No. 2.4e
0, Mismatches
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Pred. No. 2.4e+04;
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APPLICATION NUMBER: US/09/341,711

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FILE REFERENCE: 293102003220

CURRENT APPLICATION NUMBER: US/10/199,550

CURRENT FILING DATE: 2002-12-20

PRIOR APPLICATION NUMBER: US 09/963,038

PRIOR FILING DATE: 2001-09-24

NUMBER OF SEQ ID NOS: 18

SOPTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 17

LENGTH: 24
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                                                                                                                                                                                                                                                                                  RESULT 23
US-09-866-108-3580/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Porcine Adenovirus Type
US-10-199-550-17
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SEQ ID NO 36
LENGTH: 23
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Publication No. US20030099615A1
GENERAL INFORMATION:
APPLICANT: Tikoo, Suresh K.
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                            APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
PILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                   atent No.
                                                                                                                                                                                          APPLICANT: GU, Yizhong APPLICANT: JI, Yongga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 1999-09-21
APPLICATION NUMBER: PCT
FILING DATE: 1998-01-26
APPLICATION NUMBER: GB9701425.2
FILING DATE: 1997-01-24
NT FILING DATE: 2001-05-25
APPLICATION NUMBER: US 60/207,456
FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                 3580, Application US/09866108
o. US20020048800A1
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llarity 81.2%;
Conservative
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Best Local S
Matches 13
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LENGTH: 25
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                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/866;108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
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APPLICANT: JI, Yongga
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                          TILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                    PPLICANT:
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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                                                                                       FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/USO1/00666
FILING DATE: 2001-01-30
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                                                 APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
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                                   CATION NUMBER: PCT/US01/00664
                                                                                                                                                                                                                                                                                                                                                                                                             JI, Yonggang
PENN, Sharron
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81.2%;
PCT/US01/00669
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Pred. No. 2.4e+(
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LENGTH: 25
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-09-866-108-3581
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PRIOR PILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION DATE: 2000-09-27
PRIOR APPLICATION DATE: 2000-09-27
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PRIOR PPLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
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DR APPLICATION NUMBER: PCT/US01/00665

R FILING DATE: 2001-01-30

PR FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30

R FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30

DR FILING DATE: 2001-01-30

DR APPLICATION NUMBER: PCT/US01/00662
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                                                                                                                                                  APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                PILING DATE: 2000-09-27 APPLICATION NUMBER: PCT/US01/00666
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APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00665
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SHANNON, Mark
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PENN, Sharron G.
HANZEL, David K.
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Pred. No. 2
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-3582
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/866,108
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OR APPLICATION NUMBER: PCT/US01/00670
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: US 60/234,687
OR FILING DATE: 2000-09-21
OR APPLICATION NUMBER: US 60/266,860
OR FILING DATE: 2001-02-05
APPLICATION NUMBER: PCT/US01/00670 FILING DATE: 2001-01-30 APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00667
FILING DATE: 2001-01-30
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                                                                                   FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00668
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o. US20020048800A1
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INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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81.2%; Pred. No. 2.
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-3584
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                                                                                                                                                            SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 3584
                                             Query Match
Best Local
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PRIOR FILING DATE: 2000-05-26
PRIOR PPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING NUMBER: US 60/236,359
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CURRENT FILING DATE: 2001-05-25
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TYPE: DNA
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TLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
LE REFERENCE: AEOMICA-7
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APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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o. US20020048800A1
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CTGCGCCCGGATCCAG 16
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HANZEL, David K.
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Pred. No. 2.
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                              Mismatches
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                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing
SEQ ID NO 3585
LENGTH: 25
TYPE: DNA
                                                                                                      Query Match
Best Local (
                                                                                      Matches
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
APPLICANT: SHANNON, Mark
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AROMICA-7
CURRENT FAPPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
DEFIOR ETILING DATE: 2000-10-04
PRIOR ETILING DATE: 2000-10-07
PRIOR PILING DATE: 2000-10-07
PRIOR PILING DATE: 2000-10-08
PRIOR APPLICATION NUMBER: US 60/236,359
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT,
FILING DATE: 2001-01-30
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                                                                                      Conservative
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                                                                                                        Score 11.2; DB 10;
Pred. No. 2.4e+04;
                                                                                    Mismatches
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RESULT 29 US-09-866-108-3586/c

Sequence 356, Application US/09866108 Patent No. US20020048800A1 GENERAL INFORMATION:
APPLICANT: GU, Yizhong

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RESULT 30
US-09-866-108-3587/c
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SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 3586
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ORGANISM: Homo sapiens
-09-866-108-3586
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR FILING DATE: 2001-01-30
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PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
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PRIOR PILING DATE: 2001-01-30
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APPLICANT: JI, Yongga
FILE REFERENCE: AEOMICA-7
TURRENT APPLICATION NUMBER: US/09/866,108
TURRENT FILING DATE: 2001-05-25
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ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                             PPLICANT: RANK, David R.
PPLICANT: CHEN, Wensheng
PPLICANT: SHANNON, Mark
ITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
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APPLICATION NUMBER: PCT/US01/00670
PILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
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                                                                                                                                                      HANZEL, David K.
                                                                                                                                                                                                     JI, Yonggang
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L, David K.
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81.2%; Pred. No. 2
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                                                                     IN HUMAN HEART AND MUSCLE
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US-09-866-108-3587
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                                 CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR TILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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                                                                                                                                                                           FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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LENGTH: 25
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00667
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5. US20020048800A1
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PENN, Sharron G.
HANZEL, David K.
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Pred. No. 2.4e+04;
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IVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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Best Local
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CURRENT FILING DATE: 2001-05-25
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NUMBER OF SEQ ID NOS: 15752
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                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SHANNON, MARK
FITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/236,359
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                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
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APPLICATION NUMBER: US 60/207,456
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APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
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                                                                                            FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
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PENN, Sharron G.
HANZEL, David K.
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NUMBER: PCT/US01/0066
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Pred. No. 2
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RESULT 34
US-10-251-482-17
; Sequence 17, Application US/10251482
; Publication No. US20030099631A1
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Best Local Similarity
"---hes 12; Conserve
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; OTHER INFORMATION:
US-10-080-959A-5
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PRIOR FILING DATE: 2001-01-30
PRIOR PPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2001-02-05
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Matches
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Best Local Similarity
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                                                                                                                             GENERAL INFORMATION:
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                       APPLICANT: Yuichi, Hikichi
APPLICANT: Atsushi, Nishimura
TITLE OF INVENTION: NOVEL PROTEINS, THEIR PRODUCTION
FILE REFERENCE: 48712/342
CHERENT APPROXIMATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Buttner, Mark P.
TITLE OF INVENTION: Method for Detection of Stachybotrys chartarum
TITLE OF INVENTION: Samples Using Quantitative Polymerase Chain Re
FILE REFERENCE: 0001-00001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-03-29
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CURRENT FILING DATE: 2002-05-16
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                                                                                    APPLICANT: Koji,
APPLICANT: Yuici
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Stachybotrys chartarum
FEATURE:
                 CURRENT APPLICATION NUMBER: US/10/251,482
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2002-09-19
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85.7%;
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Pred. No. 2
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Pred. No. 3.
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                                                                                                                                 RESULT 36
US-10-122-434-7
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SEQ ID NO 82
LENGTH: 20
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Sequence 7, Application US/10122434
Publication No. US20030078402A1
GENERAL INFORMATION:
APPLICANT: Leon G.J. FRENKEN
APPLICANT: Cornalis P.E. VAN DER LOGT
TITLE OF INVENTION: METHOD FOR PRODUCING ANTIBODY FRAGMENTS
FILE REFERENCE: 60113/266062 - T3076(C)
                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: PCT/US98/13961
PRIOR FILING DATE: 1998-07-06
PRIOR APPLICATION NUMBER: US 08/888,982
PRIOR FILING DATE: 1997-07-07
PRIOR APPLICATION NUMBER: US 08/756,806
PRIOR FILING DATE: 1996-11-26
PRIOR FILING DATE: 1996-11-26
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PRIOR FILING DATE: 1999-07-26
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 09/143,214 PRIOR FILING DATE: 1998-08-28
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CURRENT FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: 09/506,073
PRIOR FILING DATE: 2000-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US95/07111
PRIOR TILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/250,856
PRIOR FILING DATE: 1994-05-31
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TITLE OF INVENTION:
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: primer
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                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: antisense sequence
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85.7%;
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Pred. No. 3.8e+04;
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Pred. No. 3
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Patent No. US20010029620A1
GENERAL INFORMATION:
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Publication No. US20030096239A1
GENERAL INFORMATION:
APPLICANT: Gunderson, Kevin
APPLICANT: Chee, Mark
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LENGTH: 21
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Best Local Similarity 85:7%;
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TITLE OF INVENTION: HYBRID WINTER OILSE!
FILE REFERENCE: 514412-2019
CURRENT APPLICATION UNMBER: US/99/733,151
CURRENT FILING DATE: 2000-12-08
PRIOR APPLICATION NUMBER: 09/457,037
PRIOR FILING DATE: 1999-12-08
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NUMBER OF SEQ ID NOS: 39
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CURRENT FILING DATE: 2002-04-16
PRIOR APPLICATION NUMBER: 09/487,253
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          CURRENT APPLICATION NUMBER: US/09/940,185
CURRENT FILING DATE: 2001-08-27
PRIOR APPLICATION NUMBER: US 60/227,948
PRIOR FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/228,854
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 4768
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SOFTWARE: PatentIn version 3.1
                                                                                                                                                TITLE OF INVENTION: Probes and Decoder Oligonucleotides FILE REFERENCE: A-69605-1
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ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION: primer 254
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Pred. No. 3
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Pred. No. 3.8e+04;
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; OTHER INFORMATION: PCR primer
US-10-076-157-21
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US-09-864-785-1566
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; PEATURE:
; OTHER INFORMATION: Computer Generated Probe Sequence.
US-09-940-185-964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jimenez, Alberto;
APPLICANT: Garcia, Maria Angeles Santos
TITLE OF INVENTION: Genes of purine biosynthesis from Ashbya Gossypii and the use the
TITLE OF INVENTION: in
TITLE OF INVENTION: microbial riboflavin synthesis
FILE REFERENCE: 48684DIV
CURRENT APPLICATION NUMBER: US/10/076,157
CURRENT FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: US 09/212,247
PRIOR PILING DATE: 1998-12-16
SOFTWARE: WordPerfect v. 6.1
SOFTWARE: WordPerfect v. 6.1
SOFTWARE: WordPerfect v. 6.1
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Best Local Similarity
Matches 12; Conserv
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Best Local Similarity 85.,
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Conservative
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Patent No. US20020177568A1
GENERAL INFORMATION:
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                            APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT PILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3929
SOPTWARE: Datentin vertical 10
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ORGANISM: Artificial Sequence
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NO 1566
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Joeffken, Hans Wolfgang
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85.7%;
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Search completed: June Job time: 76.7273 secs
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                                                                                                                                                               ; TYPE: RNA; ORGANISM: Artificial Sequence; FEATURE: THEN INFORMATION; Description of Artificial Sequence: Nucleic Acid US-09-864-785-1566
                                                                                                        Query Match
Best Local Similarity
Matches 11; Conserv
                                                    UGCCCCAGGCUCCAGCC 17
                                                                               TGCGCCCGGATCCAGGC 18
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           7, 2003, 11:39:16
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US-09-318-191-2
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; ORGANISM: Stachybotrys chartarum
US-09-593-012-188
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Best Local
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GENERAL INFORMATION:
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       APPLICANT: Marcel Behr
APPLICANT: Peter Small
APPLICANT: Gary Schoolnik
APPLICANT: Michael Wilson
TITLE OF INVENTION: Molecular differences between species
TITLE OF INVENTION: the M. tuberculosis complex
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HAUGLAND, Richard
APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING AND QUANTIFYING SPECIFIC FUNGI AND BACTERIA
FILE REFERENCE: HAUGLAND=1A
CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US-09/290,990
PRIOR PILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: US-09/201,773
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NUMBER OF SEQ ID NOS: 225
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US-08-624-545-50
US-08-353-476-44
US-08-460-751-19
US-08-554-137-4
US-08-663-42
US-09-168-006A-7
US-09-118-191-2
US-09-130-339-5
US-08-125-012-27
US-08-125-012-20-8
US-08-125-012-20-8
US-08-125-012-20-8
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Matches 14
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TYPE: DNA
ORGANISM: M. tuberculosis
FEATURE:
NAME/KEY: misc_feature
LOCATION: 1
                                                                                                                 GENERAL INFORMATION:
APPLICANT: Jensen.
                                                                                                      TITLE OF INVENTION:
COMPUTER READABLE
                                                                             NUMBER OF SEQUENCES: E
        STATE: New York COUNTRY: U.S.A. ZIP: 10174-6401
                                           STREET: 405 Lexington CITY: New York
                                                                  ADDRESSEE:
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CURRENT APPLICATION NUMBER: US/09/593,012
CURRENT FILING DATE: 2000-06-13
PRIOR APPLICATION NUMBER: US 09/290,990
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 225
SOFTWARE: Patentin version 3.0
SEQ ID NO 106
LENGTH: 17
TYPE: DNA
ORGANISM: Myrothecium verrucaria/roridum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: genome sequence, US-09-318-191-2
                                                                                                                       Sequence 2, Application US/08284566A Patent No. 5681725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/318,191A CURRENT FILING DATE: 1999-05-25 EARLIER APPLICATION NUMBER: 60/097,936 EARLIER FILING DATE: 1998-08-25 NUMBER OF SEQ ID NOS: 34 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: HAUGLAND, Richard
APPLICANT: VESPER, Stephen
TITLE OF INVENTION: METHOD OF IDENTIFYING
FILE REFERENCE: HAUGLAND-1A
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               56817250 No. 5681725disk of
Avenue,
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Pred. No. 1
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Pred. No.
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               5681725th America, Inc.
                                                                      PROTEINS
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US-09-506-855-28/c
; Sequence 28, Application US/09506855
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US-09-118-408-28
                                                                                                                                   RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 28 LENGTH: 24
                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 626554
                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Sheppard, Paul O.
TITLE OF INVENTION: ADIPOCYTE-SPECIFIC PROTEIN HOMOLOGS
FILE REPERENCE: 97-30
CURRENT APPLICATION NUMBER: US/09/118,408A
CURRENT FILING DATE: 1998-07-17
EARLIER APPLICATION NUMBER: 60/053,154
EARLIER FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 212 867 0298
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
              APPLICANT: Sheppard, Paul O.
APPLICANT: Lasser, Gerald W.
APPLICANT: Bishop, Paul D.
                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                   tent No. 64482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Harrington, James J.
REGISTRATION NUMBER: 38,711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 09-AUG-19
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: 1:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 36
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 867 0123
TELEPAX: 212 867 0298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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COMPUTER: II
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INHIBITORS FOR USE IN HEMOSTASIS AND
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92.9%;
                                                                                                                                                                                                                                                                    67.8%; Score 12.2; 82.4%; Pred. No. 3
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Pred. No. 2.5e+03;
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FILE OF INVENTION: IMMUNE FUNCTION
FILE REFERENCE: 99-12
CURRENT APPLICATION NUMBER: US/09/506,855
CURRENT FILING DATE: 2000-02-17
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 28
LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
FEATURE: THEOREMETERS. Oligonial Sequence
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Patent No. 60
GENERAL INFO
                                                                                        TELEFAX: (212) 751-6849
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS; single
 Query Match 66. Best Local Similarity 100 Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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MEDIUM TYPE: FLOPPY
                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/840,316
FILING DATE: 11-APR-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TILE OF
                                                                                                                                                                                                                   NAME: Richard W. Bork
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: NEW YORK
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                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                  PPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Emerson, Suzanne U., Purcell, Robert H.,
Tsarev, Sergei. A., and Robinson, Robin A.
ENTION: Recombinant Proteins Of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                FLOPPY DISK
                                                                                                                                                                         751-6849
              100.0%; Pred No.
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                                                                                                                                                                                                                                36,459
                                                                                                                                                           88:
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Pred. No. 3.1e+03
   Mismatches
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                                DB 3;
                3.8e+03
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                             Length 25;
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   Indels
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Gaps
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Use In Diagnostic Methods And Vaccines

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RESULT 9
US-08-471-971-88/c
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                                                                                    Sequence 88, Application US/08471971 Patent No. 6287759
                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                    Query Match
                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 751-6849
INFORMATION FOR SEQ ID NO: 88
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: TBATEV, SET
APPLICANT: Suzanne U.,
TITLE OF INVENTION: Re
TITLE OF INVENTION: A
TITLE OF INVENTION: UE
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                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 20:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 07/94
PILING DATE: 18-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Richard W. Bork
                         APPLICANT: Tsarev, Sergel. A., APPLICANT: Suzanne U., Purcell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: USO8, PILING DATE: 03-OCT-1994 PRIOR APPLICATION DATA;
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PRIOR APPLICATION DATA:
PRIOR APPLICATION UMBER: US08/316,765
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     TITLE OF INVENTION:
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CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: NEW YORK
STATE: NEW YORK
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FILING DATE: 03-OCT-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                      COPOLOGY:
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SYSTEM: PC-DOS/MS-DOS
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                                Sergel. A., Emerson,
U., Purcell, Robert H.
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U., Purcell, Robert H.
Recombinant Proteins Of
A Pakistani Strain Of Hepatitis E And Their
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                                                                                                                                                                                                                                                                   Score 12; ; Pred. No.
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Proteins Of
Strain Of Hepatitis
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. 3.8e+03;
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US-09-402-776-88/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US08/316,765
FILING DATE: 03-OCT-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US07/947,263
FILING DATE: 18-SEP-1992
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 751-68
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                   APPLICANT: Emerson, APPLICANT: Tearev, STITLE OF INVENTION: TITLE OF INVENTION: UMBER OF SEQUENCES:
          COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPEY DISK
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WORDPERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Richard W. Bork
REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 20
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NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 345 PAR
CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                ZIP: 10154
                                                                                                                                   CITY: NEW YORK
                                                                                                                                                   STREET:
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                                                                                                                                                                  DDRESSEE:
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                                                                                                                                                                                                                                                                                                                         8, Application US/09402776
6458562
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCGGATCCAGGC 18
                                                                                                                  NEW YORK
                                                                                                                                                  E: MORGAN & FINNEGAN
345 PARK.AVENUE
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                                                                                                 USA
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WORDPERFECT 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               66.7%;
                                                                                                                                                                                                                                     A Pakistani Strain
                                                                                                                                                                                                                                                       Recombinant
                                                                                                                                                                                                                                                                                     Suzanne U., Purcell, Robert H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US/08/471,971
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                                                                                                                                                                                                                    In Diagnostic Methods
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; Pred. No.
                                                                                                                                                                                                                                                       A., and Robinson, Robin A.
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CURRENT APPLICATION DATA:

APPLICATION NUMBER:

US/09/402,776

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RESULT 11
PCT-US93-08849A-88/c
Sequence 88, Application PC/TUS9308849A;
GENERAL INFORMATION:
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CT-US93-08849A-88
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                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US07/947,263
FILING DATE: 18-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: William S. Feiler
REGISTRATION NUMBER: 26,728
REFERENCE/DOCKET NUMBER: 2026-4032 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212) 751-68
INFORMATION FOR SEQ ID NO:
                                                                                                     TELEFAX: (212) 751-6849 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
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REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 2026-4255
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/84
PILING DATE: 11-APR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08849A
PILING DATE: 17-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY
                                             SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                    COMPUTER: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ITLE OF INVENTION: ITLE OF INVENTION:
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ADDRESSEE: MORGAN & FINNEGAN
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                              STRANDEDNESS:
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A Pakistani Strain Of Hepatitis E And Their
Use In Diagnostic Methods And Vaccines
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hes 0;
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PCT-US93-08849-88/c
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                                                                                                               RESULT 13
US-08-474-140-3/c
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                                                                                  Sequence 3, Application US/08474140 Patent No. 5721127
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Best Local (
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                                        GENERAL INFORMATION APPLICANT: DEWEEL AMORY.
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                                                                                                                                                                                                                                                                                                                                                                                                          NAME: BORK, Richard, W. REGISTRATION NUMBER: 36,459 REGISTRATION NUMBER: 20 REFERENCE/DOCKET NUMBER: 20 TELECONMUNICATION INFORMATION: TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/947,263
FILING DATE: 18-SEP-1992
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: FLOPPY DISK
TITLE OF INVENTION:
TITLE OF INVENTION:
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ADDRESSEE: MORGAN &
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                                                                                                                                                                                                                                                                                                                 TYPE: nuclead
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FILING DATE: 17-SEP-1993
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STATE: NEW YORK
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                                     AMORY, ANTOINE
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A Pakistani Strain Of Hepatitis E And Their
Use In Diagnostic Methods And Vaccines
PULLULANASE, MICROORGANISMS WHICH PRODUCE IT, PROCESSES FOR THE PREPARATION OF THIS PULLULANASE AND THE USES THEREOF
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                                                                                                                                                                                                                                              Score 12; DB 5; Length 25; Pred. No. 3.8e+03;
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, Pred. No.
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RESULT 14
US-08-477-630-3/c
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TELEFAX: (202) 293-1850
TELEX: 650 383-5605
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic grid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/08477630 Patent No. 5721128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,630
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: DEWEER
APPLICANT: AMORY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: 1ir
MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: WILLIAN BRINKS HOPER GILSON & LIONE, P.C.
STREET: 2000 K Street, N.W., Suite 200
                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
                   ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 37,136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                          CITY: Washington
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                                                                                                                                                                                                      20006
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Wilhlem F. Gadiano, Esq
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AMORY, ANTOINE
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                                                                                                            Version
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RESULT 15
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Query Match
Best Local Similarity 86. 
Matches 13; Conservative
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Patent No. 5731174
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Best Local :
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TELEX: 650 383-5605
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                          TELEX: 650 383-5605
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: I
                                                                         MOLECULE TYPE:
DESCRIPTION:
                                                                                                                                                              SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 4121-44
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 429-0625
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                     TOPOLOGY:
                                                                                                                TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                              NAME: Wilhlem F. Gadiano, REGISTRATION NUMBER: 37,1
                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: D.C.
COUNTRY: U.S.A.
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DESCRIPTION: synthetic DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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                                                                                                                                              ENGTH:
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                                                                                                                                nucleic acid
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                                                                                                                                                                                                        (202) 293-1850
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                                                                                                       linear
                                                                      nucleic acid (other);
synthetic DNA
                                                                                                                   single
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86.7%;
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                Score 11.8; DB 1;
Pred. No. 4.7e+03;
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Pred. No. 4
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     Mismatches
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                               DB 1;
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CTGCGCCCGGATCCA 15

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US-08-474-545-3/c
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APPL
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Best Local (
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APPLICANT: DEWEER, PHILIPPE
APPLICANT: AMORY, ANTOINE
TITLE OF INVENTION: PULLULANASE, MICROORGANISMS WHICH
TITLE OF INVENTION: PRODUCE IT, PROCESSES FOR THE PREPARATION OF THIS
TITLE OF INVENTION: PULLULANASE AND THE USES THEREOF
                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    equence 3, Application US/08478341 tent No. 5817498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
DESCRIPTION:
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ADDRESSEE: WILLIAN BRINKS HOFER GILSON & LIONE, P.C.
STREET: 2000 K Street, N.W., Suite 200
                                                                                                      APPLICANT: DEWEER, PHILIPPE APPLICANT: AMORY, ANTOINE TITLE OF INVENTION: PULLULAN TITLE OF INVENTION: PULLULAN TITLE OF INVENTION: PULLULAN
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ADDRESS:
WILLIAN BRINKS HOPER GILSON & LIONE, P.C
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synthetic DNA
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86.7%;
                                                                                                      PULLULANASE, MICROORGANISMS WHICH PRODUCE IT, PROCESSES FOR THE PREPARATION OF THIS PULLULANASE AND THE USES THEREOF
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Pred. No. 4.
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US-08-996-733-3/c
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TELEPAX: (202) 293-1850
TELEX: 650 383-5605
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
ILENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local 9
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                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION NUMBER: US/08/996,733
FILING DATE: 23-DEC-1997
PRIOR APPLICATION NUMBER: US 08/472,293
APPLICATION NUMBER: US 08/472,293
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COMPUTER: IBM PC compatable
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/478,341
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Wilhlem F. Gadiano, Esq.
NEGICET STREET OF THE ST
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MOLECULE TYPE:
DESCRIPTION:
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REFERENCE/DOCKET NUMBER: 41
TELECOMMUNICATION INFORMATION:
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FILING DATE: 07-JUN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 28-DEC-
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APPLICANT: AMORY, ANTOINE
TITLE OF INVENTION: PULLUI
TITLE OF INVENTION: PRODU
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CORRESPONDENCE ADDRESS:
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925 Page Mill Road
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   MBER: US 08/174,893
28-DEC-1993
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US-08-275-526C-17/c
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Best Local :
                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING YSCIEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/275,526C
FILLING DATE: 15-JUL-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
MANUE: GASTAGE WILLIAMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (650) 845-6504
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: VETTER, ROMAN TITLE OF INVENTION: XYLAND TITLE OF INVENTION: EXPRES TITLE OF INVENTION: USE TH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 33,888
REFERENCE/DOCKET NUMBER: GC446C1-US
REFERENCE/DOCKET NUMBER: GC446C1-US
REFERENCE/DOCKET NUMBER: 33,888
REGISTRATION INFORMATION:
TELEPHONE: (650) 846-7620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: BE 09201156
PILING DATE: 28-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Debra J. Glaister, Esq.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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NAME: Gadiano, wilhlem F.
REGISTRATION NUMBER: 37,136
REFERENCE/DOCKET NUMBER: 4121-49
                                                                                                                                                                                                                                                                                                       CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: BE 09300744
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6180382
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D.C.
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                                                                                                                                                                                                                                                                                                                     E: WILLIAN BRINKS HOFER GILSON & LIONE,
2000 K St., N.W., Suite 200
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LAHAYE, ANDR E
LEDOUX, PIERRE
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DETROZ, REN
ANDRE, CHRISTOPHE
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                                                                                                                                                                                                                                                                                                                                                                                                          XYLANASE DERIVED FROM A BACILLUS SPECIES, EXPRESSION VECTORS FOR SUCH XYLANASE AND OTHER PROTEINS, HOST ORGANISMS THEREFOR AND USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.6%; Score 11.8; DB 3; 86.7%; Pred. No. 4.7e+03; tive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: US-08-275-526C-17
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Matches
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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INFORMATION FOR SEQ ID NO:
                                                                                                             TELEFAX: (202) 293-0625
TELEX: 650 383 5605
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
LENGTH: 18 base pairs
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STRANDEDNESS:
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                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/076,677
FILING DATE: 12-May-1998
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION TUMBER: US 08/275,526
FILING DATE: 15-UL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 20006
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
TOPOLOGY: linear MOLEGUE TYPE: nucleic acid (synthetic oligonucleotide) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 4121-49
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 429-0625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: XYLANASE DERIVED FROM A BACILLUS SPECIES,
EXPRESSION VECTORS FOR SUCH XYLANASE AND
OTHER PROTEINS, HOST ORGANISMS THEREFOR AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington STATE: D.C.
                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                       NAME: Gadiano, Wilhlem F. REGISTRATION NUMBER: 37,136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: WILLIAN BRINKS HOFER GILSON & LIONE, P.C. STREET: 2000 K St., N.W., Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20006
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LEDOUX, PIERRE
AMORY, ANTOINE
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ANDRE, CHRISTOPHE
VETTER, ROMAN
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86.7%;
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Pred. No. 4.7e+03;
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US-09-076-677-17

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TOPOLOGY: linear

MOLECULE TYPE: nucleic acid (synthetic oligonucleotide)

BEQUENCE DESCRIPTION: SEQ ID NO: 17:

US-09-073-055-17
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                                                                                                                                                                                              TELECOMPULICATION INFORMATION:
TELEPHONE: (202) 429-0625
TELEPAX: (202) 293-0625
TELEPAX: (202) 393-5605
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pair
TYPE: nucleic acid
STRANDENESS: Single
                                                                                Query Match
Best Local S
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Patent No. 6426211
GENERAL INFORMATION:
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Best Local Similarity
                                                                Matches
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/275,526
FILING DATE: 15-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Gadiano, Wilhlem F.
REGISTLATION NUMBER: 37,136
REPERENCE/DOCKET NUMBER: 4121-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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EXPRESSION VECTORS FOR SUCH XYLANASE AND
OTHER PROTEINS, HOST ORGANISMS THEREFOR AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
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                              1 CTGCGCCCGGATCCA 15
                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 05-May-1998 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/073,055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
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                                                                Conservative
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ANDRE, CHRISTOPHE
VETTER, ROMAN
VETTER, ROMAN
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LEDOUX, PIERRE
AMORY, ANTOINE
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                                                                                65.6%; Score 11.8; DB 4; Length 18; B6.7%; Pred. No. 4.7e+03;
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Pred. No. 4.7e+03;
                                                                Mismatches
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                                                                Indels
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                                                                Gaps
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RESULT 23
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US-08-425-299A-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPAX: (617) 227-5941
INFORMATION FOR SEQ ID NO: 13
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDENESS: single
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Patent No.
                                                                                                                                                                                                                                        Sequence 25, Application US/08480173A
Patent No. 6072049
GENERAL INFERMATION:
APPLICANT: Thoma, Hans A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII(text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/425,299A
FILING DATE: 20-APR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                              ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: HMI-014
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 227-7400
                                                                                                                                                                                TITLE OF INVENTION: HEPATITIS NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE MEDIUM TYPE: FI
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ADDRESSEE: LAHIVE & COCKFIELD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Peters, Jean-Michael
TITLE OF INVENTION: Assay and Reagents for Detecti
TITLE OF INVENTION: of Ubiquitin-Dependent Degrada
TITLE OF INVENTION: Cell Cycle Regulatory Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kirschner, Marc
APPLICANT: King, Randall W
APPLICANT: Peters, Jean-Mic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ELECUTATION (617)
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                             STATE:
                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 65.6%;
Local Similarity 86.7%;
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                                                                                                                              CITY: Minneapolis
                                                                                                                                              ADDRESSEE: Popovich & Wiles, P.A.
STREET: 80 S. 8th Street, Suite 1902
                                                                                                                                                                                                                                                                                                                                                                                                                              3 GCGCCCGGATCCAGG 17
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Pred. No. 4.7e
0; Mismatches
                                                                                                                                                                                                                          B SURFACE ANTIGEN VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8; DB 1;
4.7e+03;
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RESULT 24
US-08-480-173A-26
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         ATTORNEY/AGENT INFORMATION:
NAME: Popovich, Thomas E
REGISTRATION NUMBER: 30,099
REFERENCE/DOCKET NUMBER: MED1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-334-8991
                                                                                                                                                                                                                                                                                                                                                                                                  atent No.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOPTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                     PPLICANT: Thoma, Hans A.
                                                                                    CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                           STREET: 80 S. 8th CITY: Minneapolis
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REFERENCE/DOCKET NUMBER: MED1003USD4
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                                                                                                                                                                                                                                             COUNTRY:
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5. 6072049
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pedness: double
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80 S. 8th Street, Suite 1902
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612-334-8994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.6%;
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single-stranded "sticky end""
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                                         MED1003USD4
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Pred. No. 4.7e+03;
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                                                                                                                                                         Version #1.30
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Matches
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GENERAL INFORMATION:
                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/00
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy
                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Popovich, Thomas E
REGISTRATION NUMBER: 30,0
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                                                                                                        NOLECULE TYPE:
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                        NAME/KEY:
LOCATION:
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SOPTWARE: PatentIn Polo-
                              OTHER INFORMATION: OTHER INFORMATION:
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CITY: Minneapolis
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OTHER INFORMATION:
NAME/KEY: misc_feature
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13; Conserv
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EDNESS: double
                                                                                                                                             nucleic acid
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80 S. 8th Street, Suite 1902
                                                                                                                                                                                                                  612-334-8994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thoma, Hans A
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                                                                      misc_feature
                                                                                                                        linear
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                                                                                                      DNA (synthetic)
                                                                                                                                      double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HEPATITIS B SURFACE ANTIGEN VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.6%; Score 11.8;
86.7%; Pred. No. 4.
                           /note= "Nucleotides 1-4 single-stranded "sticky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Adapter sequence results
from oligonucleotide duplex formation with nucleotides 5-24
SEQ ID NO: 25"
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                                                                                                                                                                                                                                                                                                                                                         US/08/484,408A
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                                                                                        Query Match
                                                            Matches
                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 612-334-8994
INFORMATION FOR SEQ ID NO:
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tent No. 6117653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: PATENTIA RE-BASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,408A
PILING DATE: 07-UN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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OTHER INFORMATION:
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OTHER INFORMATION:
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LOCATION: 5..24
OTHER INFORMATION:
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                                                                                                                                                                                                                                  LOCATION: 1..4
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                LENGTH: 24 base pairs TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Popovich, Thomas E
REGISTRATION NUMBER: 30,099
REFERENCE/DOCKET NUMBER: MED1003USD4
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                               TGCGCCCGGATCCAG 16
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80 S. 8th Street, Suite 1902
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86.7%;
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                                                                           65.6%;
                                                                                                                                                                                                                                 /note= "Nucleotides 1-4 form a single-stranded "sticky end""
                                                                                                                                      /note= "Adapter sequence results from oligonucleotide duplex formation with nucleotides SEQ ID NO: 25"
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Pred. No. 4.7e+03
0; Mismatches
                                                                           Score 11.8; DB 3;
Pred. No. 4.7e+03;
                                                            Mismatches
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GENERAL INFO

Oeystein, Fodstad Hovig, Eivind Engebraaten, Olav Maelandsmo, Gunhild H.

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                                                                                                                                                                        Matches
                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 661-80 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/(
FILING DATE: 13-OCT-199-)
CLASSIPICATION: 800
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: FRANSEN, LUCIA; DEVOS, K. APPLICANT: ANDRE; VAN HEUVERSWYN, HTITLE OF INVENTION: NEW POLYPEPTIDES TITLE OF INVENTION: CODING TO THE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/BI
FILING DATE: 28-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 92.401
FILING DATE: 30-APR-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
ADDRESSEE: BIERMAN AND MUSERLIAN
                                                                                                                                                                                                                                                                             ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: CHARLES A. MUSERLIAN
REGISTRATION NUMBER: 19,68
                                                                                                                                                                                                                                                                                                                           ropology:
                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                       PE: nucleic acid
RANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                                                                                       25 base pairs
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CODING FOR THEM, AND THEIR USE IN THE FIELD OF TUMOR THERAPY O
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Pred. No. 4
                                                                                                                                                                        Mismatches
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HUGO
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                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
          COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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ADDRESSEE: HALE AND DORR LLP
STREET: 60 State Street
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REFERENÇE/DOCKET NUMBER: HYZ-039CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rel
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MEDIUM TYPE: Floppy
                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                            ITTLE OF INVENTION: CAPL SPECIFIC OLIGONUCLEOTIDES AND
                                                                                                                           ADUAL STREET: STREET: Boston
                                                                                                                                                                                                              TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TUMBER OF SEQUENCES:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Polocom
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/602,036A FILING DATE: 16-FEB-1996
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nucleic acid
DEDNESS: single
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APPLICATION DATA:
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                                                                                                                                                            60 State Street
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                                                                                                              United States of America
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Maelandsmo, Gunhild H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            64.4%; Score 11.6; 77.8%; Pred. No. 5.
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APPLICANT: Oeyste
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                           REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
                                                                                                                                                                                                             APPLICATION NUMBER: US/01
FILING DATE: 16-FEB-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: HY TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
MOLECULE TYPE:
                                                            SEQUENCE CHARACTERISTICS LENGTH: 22 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: CAPL SPECIFIC OLIGONUCLEOTIDES AND
TITLE OF INVENTION: METHODS OF INHIBITING METASTATIC CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                 TOPOLOGY: 1:
                                                                                                                                                                              NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
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TOPOLOGY: lir
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Maelandsmo, Gunhild H.
                                                                                                               (617)
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Pred. No. 5.8e+03;
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; HYPOTHETICAL:
; ANTI-SENSE: N
US-08-602-036A-3
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US-08-602-036A-4/c
                            RESULT 32
US-08-502-374A-3/c
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Best Local Similarity 77.0
Matches 14; Conservative
Sequence 3, Application US/08502374A Patent No. 5872007
                                                                                                                                                                  Best
                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 526-5000 NFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/602,036A
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-039CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
STREET: 60 State Street
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                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Kerner, Ann-Lo
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                   ENGTH:
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                                                                                                           1 CTGCGCCCGGATCCAGGC 18
                                                                                                                                               Similarity 77.8
14; Conservative
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                                                                                       CTGCTCCCAGATCCTGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INVENTION: METHODS OF INHIBITING METASTATIC CANCER
                                                                                                                                                                                                                                                                                                                   22 base pairs
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Hovig, Eivind
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Maelandsmo, Gunhild H.
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                                                                                                                                                             Score 11.6; DB 1;
Pred. No. 5.8e+03;
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Pred. No. 5.8e+03
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US-08-502-374A-3
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                            tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/502,374A
FILING DATE: 14-Jul-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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HYPOTHETICAL:
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                                                                                                                                                  APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: CAPL-SPECIFIC OLIGONUCLEOTIDES AND
TITLE OF INVENTION: METHODS OF INHIBITING METASTATIC CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE:
                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:
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                              COUNTRY:
                                                  CITY: Boston
STATE: Massa
                                                                                  STREET:
                                                                                                    ADDRESSEE:
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60 State Street
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Maelandsmo; Gunhild H.
Agrawal, Sudhir
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Hovig, Eivind
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Maelandsmo, Gunhild H.
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Pred. No. 5.8e+03
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Best Local Similarity
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             TELEFAX: (617) 526-5000 INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/08/642,407A
FILING DATE: 03-May-1996
CLASSIFICATION: 514
CTARSIFICATION: 514
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APPLICATION NUMBER: US/08/502,374A
FILING DATE: 14-Jul-1995
CLASSIFICATION: 514
                                                 ATTORNEY/AGENT INFORMATION:
NAME: KETNEY, ANN-LOUISE
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
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SEQUENCE CHARACTERISTICS
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\PPLICANT: Maelandsmo, Gunhild H.
\PPLICANT: Agrawal, Sudhir
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TITLE OF INVENTION: CAPL SPECIFIC OLIGONUCLEOTIDES AND
TITLE OF INVENTION: METHODS OF INHIBITING METASTATIC CANCER,
UMBER OF SEQUENCES: 23
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REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
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617-526-5000
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Pred. No. 5.8e+03;
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                                                       Matches
                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 03-May-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                   NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
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                                                                                                                                                 TOPOLOGY: 11
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TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                TYPE: I
                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                                 STRANDEDNESS:
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ZIP: 02109
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14; Conservative
                                                                    Similarity
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nucleic acid
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Maelandsmo, Gunhild H.
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77.8%;
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                                                                   Score 11.6; DB 2
Pred. No. 5.8e+03
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Pred. No. 5.8e+03;
                                                       Mismatches
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US-08-585-595-35/c
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Patent No. 6271029
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 17
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ANTISENSE MODULATION OF CYTOHESIN-2 EXPRESSION PILE REPERENCE: RTS-0096
CURRENT APPLICATION NUMBER: US/09/428,583
CURRENT FILING DATE: 1999-10-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                       NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,595
PILING DATE: 16-JAN-1996
CLASSIPICATION: 435
                                                                                                                                                     NAME: AUERBACH, JEFFREY I.
REGISTRATION INVMER: 32600
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 383-7451
                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
PILING DATE: 11-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                               TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER OF SEQUENCES:
                                                                                                   EQUENCE CHARACTERISTICS:
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nes 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1299 PEN CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: HOWREY & SIMON
STREET: 1299 PENNSYLVANIA AVENUE, N.W.
                                                 STRANDEDNESS:
                CULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35, Application US/08585595
5. 5795761
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                                                            : 22 base pairs nucleic acid
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VVENTION: IMPROVED METHODS FOR PRODUCING VITAMIN C
                                                                                                                                        (202)
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                                 linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                 single
                DNA (genomic)
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                                                                                                                                                                                                                                                                                    US 08/584,019
                                                                                                                                                                                              6137-0014 CIP
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Pred. No. 7.
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                                  US-08-244-122-25/c
                                                    RESULT 39
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// MOLECULE TYPE: Synthetic DNA
US-09-142-623-20
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Matches
Sequence 25, App. -- No. 5866777
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                                                                                                                                                                                                                                                                                                                   TELEFAX: 202-721-8250
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/: FILING DATE: September 10, PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: KOJÍ YANAI et al.
TITLE OF INVENTION: -FRUCTOFURANOSIDASE AND ITS GENE, METHOD OF
TITLE OF INVENTION: ISOLATING -FRUCTOFURANOSIDASE GENE, SYSTEM
TITLE OF INVENTION: -FRUCTOFURANOSIDASE, AND -FRUCTOFURANOSIDA
                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 40,949
REFERENCE/DOCKET NUMBER: 98-0989*/LC(WMC)/144
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-721-8200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM:
                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: D.C.
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Local Similarity 92.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICATION NUMBER:
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                                                                                                                                                             12; Conservative
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                                                                                                                                                                                                                                                                                        Nucleic acid
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                                                                                                                                                                                                                                                                                                        22 base pairs
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2033 K Street, N.W., Suite 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORYNEBACTERIUM SP
                                                                                                                                                                                                                                                          Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IBM Compatible
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Pred. No. 7.1e
0; Mismatches
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Pred. No. 7
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Patent No. 5000. Timor

Application US/08244122

APPLICANT:

Sijmons, Peter J

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US-08-679-645-531
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INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER: PCT/EP92/02559
PILING DATE: 02-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC/XT/AT or COPERATING SYSTEM: DOS
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage

MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage

COMPUTER: DISK PC/XT/AT or compatibles
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 24 BASE PAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 30086
REFERENCE/DOCKET NUMBER: U-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 708-1800
                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 10-JAN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                              Local
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FILING DATE: 20-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/01
FILING DATE: 19-MAY-1994
CLASSIFICATION: 800
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                                                                                                                                                 INFORMATION:
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New York
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Similarity 92.3%;
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Skokut, Trom-
Young, Scott A.
Folkerts, Otto
Marlo, Donald J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (212) 246-8959
                                                                                                                Edington, Brent E.
                                                                                  McSwiggen, James A.
Merlo, Patricia Ann Owens
                                                                                                                                 Zwick, Michael G.
                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        LINEAR
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                                                                                                                                                                                                                                                                                                                                                                                           YES
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                                                                       Lining
                                                        Thomas A.
COMPOSITION AND METHODS FOR
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Pred. No. 7.1e+03;
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Search completed: June 7, 2003, 09:45:39
Job time : 47.9636 secs
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                                                                                                                                                                                                                                                          TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/001,135
FILING DATE: July 13, 1995
APPLICATION NUMBER: 08/300,726
FILING DATE: September 2, 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                 TOPOLOGY: 11
                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                        TELEPHONE: (2)
                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/
FILING DATE: July 12, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Con
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                       TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 90071-2066
                                                                                                                                       Local
                                                                                                                    ch 62.2%;
l Similarity 68.8%;
ll; Conservative
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                                                           16
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                                                                                                                                                                                                                                                                                                                                                     219/247
                                                                                                                                  Score 11.2; DB 4;
Pred. No. 8.9e+03;
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                                                                                                                      Mismatches
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